

CONFORMATION ANALYSIS OF SOME CYCLIC COMPOUNDS
BY
MOLECULAR MECHANICS AND X-RAY DIFFRACTION

A Thesis presented for
the degree of Doctor of Philosophy
in the
Faculty of Science
of the
University of Glasgow
by
Michael Henry Patrick Guy

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ERRATA

Page Line

- 15 22 Delete "of the structure invariants".
Insert "arising from the Σ_1 relationships".
- 17 4 Delete "second Fourier synthesis".
Insert "structure factor calculation".
- 62 16 Add footnote "\$ For the Σ_1 relationships only".
- 98 25 Insert "approximate" before "C₂".
- 138 7 Add footnote "\$ For the Σ_1 relationships only".
- 322 8 Insert "energy" after "minimum".
-

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SUMMARY

This thesis is composed of two parts. In part I the x-ray structural analyses of two sesquiterpenoids and two derivatives of 2-naphthol are reported, while in part II the results from molecular mechanics calculations are used to investigate the conformational processes occurring in a series of cyclo-octa-1,5-dienes and to interpret some of the stereochemical peculiarities of germacrane based sesquiterpenoids.

The crystal structure analysis of the germacranolide, costunolide, was undertaken to determine its structure and stereochemistry. An incomplete structure was available at the commencement of this study, derived partially from x-ray diffraction measurements using Mo - K α radiation and partially from synthetic and spectral observations. The diffraction data were recollected using Cu - K α radiation in order to obtain more accurate measurements of the weaker reflexions. This technique was completely successful and allowed the structure and stereochemistry to be determined.

Crystals of the guianolide, mikanokryptin, were supplied by Professor W. Hertz who, on the basis of UV, IR, and NMR spectra, had elucidated most of the stereochemistry. The configurations at C(1), C(6), and C(10) remained uncertain and the x-ray analysis removed the ambiguities at these chiral atoms. In addition, molecular mechanics calculations were carried out for the isolated 1-methylenecycloheptane system of mikanokryptin and these allowed observations to be made pertaining to the stereochemistry of this and other guianolides.

Structure analyses of 1-methyl- and 1-isopropyl-2-naphthyl acetate were carried out as part of a series of

investigations, by Dr. J. Carnduff, into the autoxidation of 2-naphthols. It was anticipated that the isopropyl derivative would have been substantially more strained than the 1-methyl compound because of differences in the rates of autoxidation of the corresponding 2-naphthols. The observed structures did not support this view.

Molecular mechanics calculations for syn-3,7-dibromo-cis, cis-cyclo-octa-1,5-diene found the twist-boat conformation as the global minimum, the geometry of which was very similar to that found in an earlier x-ray analysis. The complete pseudorotational cycle was also investigated using this technique. A similar study was carried out for the anti-3,7-dibromo isomer. The conformational processes in the parent diene, cis,cis-cyclo-octa-1,5-diene, were the subject of a variable temperature NMR investigation by Professor F. A. L. Anet and molecular mechanics calculations were used to interpret these spectra and to reproduce the minimum energy geometry found by electron diffraction measurements. The anti-3,7-dimethyl derivative has recently been synthesized by Professor P. Heimbach. He kindly supplied a sample of his product for variable temperature NMR analysis and molecular mechanics calculations have been performed with a view to interpreting the spectra.

The final chapter reports a series of calculations on the stereochemistry of germacranolides. The force field used was able to reproduce the geometry of costunolide, which had been the subject of an x-ray analysis described earlier in this thesis. Subsequent calculations on the two frequently occurring conformations of the cyclodecadiene system in germacranolides have been used to investigate the stabilities of the various modes of

lactone-cyclodecadiene fusion apparent in these sesquiterpenoids and to correlate the conformation of the lactone function with both the fusion and the Cotton effect at 260nm.

PART I

X-RAY DIFFRACTION

CHAPTER ONE

Some aspects of crystallography

1.1 Historical

Modern crystallography is based on the suggestion made by Max von Laue (1912) that the regularly spaced repeating units of a crystal might act as a three-dimensional grating for x-rays. Following the experimental verification of this hypothesis by Friedrich and Knipping, the use of monochromatic radiation and the advent and improvement of the rotating crystal method made it possible to both index and measure the intensity of individual reflexions. The subsequent development of crystallography has been concerned with the sophistication of these basic techniques and the solution of the phase problem.

The phase problem arose because of our inability to refract x-rays and thus obtain an image of the molecular structure.

The phase amplitudes can be derived fairly easily from the intensities but mathematical reproduction of the image required information about the phases. From 1930 onwards a great deal of effort was spent on this task and solutions applicable to certain limited situations were found, but no general approach has been available until recent years.

For many compounds it is now possible to derive the phase relationships directly from the intensity data using a combination of inequality and statistical methods.

The remainder of this chapter is devoted to a more detailed discussion of the specific techniques which have been used in the structure analysis of the compounds which form the subject of the x-ray crystallographic

section of this thesis.

1.2 Corrections to the measured intensities

(a) Lorentz - Polarisation

Lorentz

The time required for a reciprocal lattice point to pass through the sphere of reflexion is not constant but depends on both the Bragg angle and the method of data collection.

For the four circle diffractometer with normal beam geometry the Lorentz factor, L, is given by:

$$L = 1/\sin 2\theta \quad (1)$$

Polarisation

Partial polarisation of the x-ray beam occurs on reflexion and hence the intensity of the reflected beam is corrected by a factor, p, where:

$$p = \frac{1}{2}(1 + \cos^2 2\theta) \quad (2)$$

A more complex expression is used when the incident beam is itself partially polarised.

The combined factor, L_p, for a four circle diffractometer is therefore:

$$L_p = \frac{1 + \cos^2 2\theta}{2 \sin 2\theta} \quad (3)$$

(b) Counting Loss

Automatic data collection instruments use scintillation or proportional counters to measure intensities. When very strong reflexions are present these devices become overloaded and underestimate the true count. One method of overcoming this is to remeasure the low order reflexions at reduced beam intensity and thus obtain a conversion factor for the weaker reflexions which may then be applied to the intense reflexions.

The two intensities for each reflexion may be considered to be related by the following expression:

$$J = P_1 I + P_2 I^2 \quad (4)$$

where J = intensity at reduced setting

I = intensity at normal setting

and P_1, P_2 are constants to be determined.

Each measurement of J has a standard deviation σJ and can be assigned a weight W , such that:

$$W = 1/(\sigma J)^2$$

The difference Δ between the counts for a given reflexion under the two sets of conditions is:

$$\Delta = J - (P_1 I + P_2 I^2) \quad (5)$$

and if $M = \sum W \Delta^2$ then the stationary value of M occurs when:

$$\frac{\partial M}{\partial P_1} = \frac{\partial M}{\partial P_2} = 0 \quad (6)$$

Thus differentiation of equation (5) with respect to P_1 and then P_2 yields two simultaneous equations which can

be solved for the conditions set by equation (6) to yield \underline{P}_1 and \underline{P}_2 . Resubstitution of these constants in equation (5) with the appropriate value of \underline{J} will then allow a corrected \underline{I} to be calculated for each strong reflexion.

1.3 Structure Factors

Structure factors provide the link between the observed data and the calculated electron density distribution. They are defined by two quantities, an amplitude $|F|$ and a phase angle α . $|F|$ is directly related to the measured intensity, \underline{I} , by:

$$|F| \propto I^{\frac{1}{2}} \quad (7)$$

It can be shown that for any reflexion, \underline{hkl} , the structure factor $F(hkl)$ is given by:

$$F(hkl) = \sum_{j=1}^N f_j \exp 2 \pi i (hx_j + ky_j + lz_j) \quad (8)$$

Where atom \underline{j} is situated at a point defined by the fractional atomic coordinates (x_j, y_j, z_j) and has an atomic scattering factor \underline{f}_j .

Equation (8) may also be expressed in terms of its real and imaginary parts, \underline{A} and \underline{B} , respectively, so that:

$$F = A + iB \quad (9)$$

where

$$A = \sum_{j=1}^N f_j \cos 2 \pi (hx_j + ky_j + lz_j)$$

and

$$B = \sum_{j=1}^N f_j \sin 2 \pi (hx_j + ky_j + lz_j)$$

The amplitude and phase of $F(hkl)$ are then given by:

$$|F(hkl)| = (A^2 + B^2)^{\frac{1}{2}} \quad (10)$$

$$\text{and } \alpha(hkl) = \tan^{-1}(B/A)$$

The term f_j in equation (8) and subsequent equations is a function of atom type and $\sin(\theta/\lambda)$. It is expressed in terms of the appropriate number of electrons located at the centre of the atom (e.g. C = 12). Its value diminishes with $\sin(\theta/\lambda)$ because the finite size of the electron cloud produces phase differences between waves scattered from distinct points within its volume.

Thermal motion causes the electron density to distribute itself over a larger volume with increasing temperature. The resulting reduction in atomic scattering power is greater for higher values of $\sin(\theta/\lambda)$.

For isotropic vibration the scattering factor is given by:

$$f_j = f_0 \exp[-B(\sin^2 \theta)/\lambda^2] \quad (11)$$

where f_0 = the scattering power of stationary atom

$B = 8\pi^2 \bar{U}^2$ the Debye-Waller factor

\bar{U}^2 = the mean square amplitude of the atomic displacement at right angles to the reflecting plane.

and for anisotropic vibration:

$$f_j = f_0 \exp \left[- (B_{11}^2 h^2 a^{*2} + B_{22}^2 k^2 b^{*2} + B_{33}^2 l^2 c^{*2} + 2B_{12}^{**} hka^*b + 2B_{13}^{**} hla^*c + 2B_{23}^{**} klb^*c) / 4 \right] \quad (12)$$

Where B_{11} , B_{22} etc. are the anisotropic components of the term \underline{B} in equation (11) and a^* , b^* , c^* , are the lengths of the reciprocal cell edges.

Debye-Waller factors are sometimes expressed in terms of \underline{U} where $\underline{U} = \overline{U}^2 = B/8 \pi^2 = B/78.96$

1.4 Structure factors and the electron density distribution

Any quantity such as the electron density $\rho(xyz)$ which varies periodically may be analysed into a number of separate basic components which when superimposed represent that quantity. This method of analysis is due to the French physicist J.B. Fourier and the three dimensional summation for electron density gives:

$$\rho(xyz) = \frac{1}{V} \sum_{h=-\infty}^{+\infty} \sum_{k=-\infty}^{+\infty} \sum_{l=-\infty}^{+\infty} F(hkl) \exp \left[-2\pi i(hx+ky+lz) \right] \quad (13)$$

Thus if $F(hkl)$ is known then equation (13) will reproduce the electronic distribution in the crystal. Unfortunately, only $|F(hkl)|$ is immediately available from the intensity measurements and the phase angle α must still be determined. For centrosymmetric space groups α can only take on values of 0 or π but for non-centrosymmetric cells it may assume a whole range of magnitudes and it is more convenient to rewrite equation (13) as:

$$\rho(xyz) = \sum_{h=-\infty}^{+\infty} \sum_{k=-\infty}^{+\infty} \sum_{l=-\infty}^{+\infty} \frac{|F(hkl)|}{V} \exp \left[2\pi i(hx+ky+lz) - \alpha(hkl) \right] \quad (14)$$

Where $\alpha(hkl)$ is the phase angle appropriate to $|F(hkl)|$.

There are a number of methods for obtaining the elusive $\alpha(hkl)$ and the following section describes the general approach made to this problem by contemporary crystallography.

1.5 Direct Methods of phase determination

Modified structure factors

Direct methods rely on comparing the relative values of related intensities, but because scattering power is not independent of the Bragg angle, this comparison is less straightforward than might at first appear. High angle reflexions are considerably diminished by the fall-off in scattering ability and "reflexions" may be unobserved. A correction can be applied using a unitary structure factor $\underline{U}(h)$ instead of $\underline{F}(h)$ so that:

$$U(h) = F(h) / \sum_{j=1}^N f_j \quad (15)$$

Where the summation is carried out over the \underline{N} atoms in the molecule and \underline{f}_j is the scattering factor for atom \underline{j} compensating for both theta fall-off and thermal effects as discussed in 1.3

This equation (15) expresses each structure factor as a proportion of its maximum possible value for the particular magnitude of theta at which it occurs. Thus all unitary structure factors must lie within the range ± 1 .

The extrema corresponding to those instances where all the atoms scatter in phase and hence restricting possible atomic sites. Values of ± 1 are rarely observed but greater constraints are placed on atomic location with increasingly closer approach to unity. However, the average value of \underline{U} is dependent on the number of atoms, \underline{N} , in the molecule and for reasonably sized molecules there may be so few large \underline{U} s that an insufficient number of relationships can be established. Equation(16) gives the root mean square value of \underline{U} , \underline{U}_{rms} , in terms of the total scattering power represented by each atom:

$$\underline{U}_{rms} = \left[\sum_1^N (n_i^2) \right]^{\frac{1}{2}} \quad (16)$$

where $n_i = f_i / \sum_j f_j$

A normalised structure factor, $\underline{E}(h)$, may be obtained by relating each structure factor to \underline{U}_{rms} and thus the difficulty arising with increasing \underline{N} can be overcome. This gives:

$$\underline{E}(h)^2 = |F(h)|^2 / \sum_{i=1}^N f_i^2 \quad (17)$$

Under some circumstances space group symmetry produces special sets of reflexions which are abnormally higher (or lower) than the calculated average and the \underline{E} values for these must be modified by a factor $\underline{\epsilon}$ thus:

$$\underline{E}(h)^2 = |F(h)|^2 / \epsilon \sum_{i=1}^N f_i^2 \quad (18)$$

The distribution of $|\underline{E}|$ gives an indication of the centricity or otherwise of the space group involved.

Inequality relationships

The unitary and normalised structure factors may be expressed in exactly the same way as ordinary structure factors:

$$U(h) = 2 \sum_i^{N/2} n_i \cos 2 \pi i h x_i \quad (19)$$

for a centrosymmetric space group.

Harker and Kasper found that they were able to represent equation (19) in terms of a classical Cauchy inequality and obtained the relationship:

$$|U(h)|^2 \leq \frac{1}{2} [1 + U(2h)] \quad (20)$$

This important equation allows the structure factor of an \underline{h} (or hkl) reflexion to be compared with a $\underline{2h}$ (or $2h \ 2k \ 2l$) and may enable a phase determination to be made. Similar inequalities can be derived for other symmetry elements. Inequality relationships on their own are, however, of limited value, as the number of atoms increases, since definite determination requires large values of \underline{U} .

Probability methods

Inequalities represent the limiting case of probability methods which in general produce phases which are less reliable. Much of the early work to combine probability theory with inequalities is due to (a) Sayre and (b) Cochran and Woolfson.

The basic relationships are:

$$(a) \quad S(F_{hkl}) \approx S(F_{h'k'l'}) \cdot S(F_{h-h' \ k-k' \ l-l'}) \quad (21)$$

$$(b) \quad P(E) = \frac{1}{2} + \frac{1}{2} \tanh \left(\left| E_{hkl} E_{h'k'l'} E_{h-h' \ k-k' \ l-l'} \right| \cdot N^{-\frac{1}{2}} \right)$$

In equation (21a) $\underline{S}(F_{hkl})$ means the sign of the structure amplitude $|F_{hkl}|$. $\underline{S}(F_{hkl})$ may be replaced by $\theta(F_{hkl})$, thus allowing the Sayre expression to be used for non-centrosymmetric space groups. The starting magnitudes and/or signs must obviously be reliable in order to yield reasonable atomic co-ordinates.

Equation 21b) expresses in terms of \underline{E} values the probability, $P(\underline{E})$, that for a given hkl and related pair of reflexions, the relationship in equation(21a) is true.

Practical phase determination

The best known application of equations (21) is probably the symbolic addition method of the Karles which is the basis of the program MULTAN (Main and Woolfson).

The principal subprograms are:

(a) The SIGMA 2 program which derives $|\underline{E}|$ from the observed data and then uses the largest values to set up all the triple relationships of the form expressed by equation (33a).

(b) CONVERGE uses the supplied space group information to decide which parity groups are required to define the origin and then calculates the probabilities for the signs of the structure invariants. By a process of elimination the program should eventually be able to converge on the best starting set.

(c) FASTAN uses this starting set to determine the phases of other high value $|\underline{E}|$ s. As the non-origin

defining reflexions in the starting set may take on more than one value several complete sets of phases may be generated from the [2 phase relationships. Each reflexion with $|E|$ above a specified value will have its phase calculated and this phase may be assigned a weight which is related to the probability of it being correct. FASTAN, by using a weighted version of the tangent formula puts less emphasis on the poorly determined phases in subsequent phase development and thus allows rapid convergence on the most consistent complete set from each starting combination. A figure of merit based on the degree of internal consistency is given to each complete phase set.

(d) FOURIER computes an E map using the set showing the best figure of merit from which it should be possible to recognise atomic sites.

MULTAN does not always give a correct solution. This may be due to a number of reasons, for example, poorly diffracting crystals, unreliable data, a large number of atoms in the asymmetric unit, or the particular space group concerned.

1.6 Fourier Synthesis

Phases obtained by direct or other methods are only approximate and must be further refined to yield a more accurate structure. This is achieved by combining the calculated phases with the observed amplitudes in a summation such as that represented by equation (14).

The resulting electron density distribution will not be very accurate, but it should give a better indication of atomic sites. These new atom locations can then be used in a second Fourier synthesis to work backwards and yield a calculated set of structure factors (F_c) and more accurate phases. The process may be continued in this manner until the atomic parameters cease to change significantly.

A further Fourier summation can then be carried out based on the quantity ($F_o - F_c$). This will reveal any non-hydrogen atoms inadvertently missed during the F_o synthesis and often indicates the position of many of the hydrogen atoms.

1.7 Least squares refinement

Fourier summation as described above will contain termination of series errors, since the amount of data is finite. The reasonably correct atomic locations available from the Fourier synthesis may now be further refined by the method of least squares. This method is based on the principle that the best fit, between observed data and related functional equations, is obtained when the sum of the squares of the differences between observed and calculated values is a minimum. The function, D , is the one most commonly used:

$$D = \sum_{hkl} W(|F_o| - |F_c|)^2 = \sum_{hkl} W\Delta^2 \quad (22)$$

Where W is a measure of the accuracy of each observation.

If the variables affecting $|F_c|$ are represented by $P_1, P_2, \dots, P_j \dots P_n$ then the stationary value of equation (22) is given by:

$$\frac{\partial D}{\partial P_j} = 0 = \sum_{hkl} W \Delta \frac{\partial |F_c|}{\partial P_j} \quad (j = 1, n) \quad (23)$$

These n equations (23) have a solution which will give the best values of the parameters P_j . The large number of equations involved for an average sized structure cannot be solved easily. However, values for the P_j are already available from the F_0 summation and at this stage only small changes, $\underline{\epsilon}$, should be required. The total change $\underline{\Delta}$ may be expanded as a truncated Taylor series:

$$\Delta(P + \underline{\epsilon}) = \Delta P - \sum_{i=1}^n \epsilon_i \frac{\partial |F_c|}{\partial P_i} \quad (24)$$

Where subscripted/non-subscripted variables refer to individual/total changes and parameters.

Combination of equations (23) and (24) gives a set of normal equations:

$$\sum_{i=1}^n \left\{ \sum_{hkl} W \frac{\partial |F_c|}{\partial P_i} \cdot \frac{\partial |F_c|}{\partial P_j} \right\} \epsilon_i = \sum_{hkl} W \Delta \frac{\partial |F_c|}{\partial P_j}$$

The normal equations may be written as:

$$\sum_i a_{ij} \epsilon_i = b_j \quad (j = 1, 2, \dots, n) \quad (25)$$

Solution of these sets of normal equations will yield the shifts $\underline{\epsilon}_i$ and the calculation may be repeated to convergence by using the new P_j from one cycle as the starting point for the following one.

Least squares refinement can be carried out using either block diagonal or full matrix techniques. The saving in computer time and storage for the former must be offset against the possibility that the off-diagonal relationships are not small.

The errors in the final set of atomic co-ordinates may be estimated from the following expression:

$$\sigma(P_i) = \left[(a^{-1})_{ii} \frac{\sum W \Delta^2}{m - n} \right]^{\frac{1}{2}} \quad (26)$$

Where $\sigma(P_i)$ is the error in parameter P_i .

$(a^{-1})_{ii}$ is the inverse of a matrix of the form represented by equation (25).

m is the number of observations.

n is the number of parameters.

The quantity W should be given by:

$$W(hkl) = \frac{1}{\sigma^2(hkl)}$$

Where $\sigma^2(hkl)$ is the esd of the amplitude $|F_o(hkl)|$ and this could be determined by repeated observations of each reflexion. This is clearly impracticable for a large body of data and an empirical weighting scheme is usually employed instead. Such schemes are designed to keep the function $W(|F_o| - |F_c|)^2$ constant when averaged over batches of data.

The quality of fit obtained in a least squares refinement may be expressed in terms of a weighted residual, R :

$$R = \frac{\sum w(|F_o| - |F_c|)^2}{\sum w |F_o|^2} \quad (27)$$

CHAPTER TWO

X-ray diffraction studies of the sesquiterpenoids
costunolide and mikanokryptin

2.1 Foreword

Sesquiterpenoids are C₁₅ compounds which may be considered (1,2) to be built up from three isoprenoid units. A number of different cyclisations of the C₁₅ precursor can occur giving rise to the wide diversity of structural types currently known. The most common carbon skeletons (3) are illustrated in Fig. 2.1.

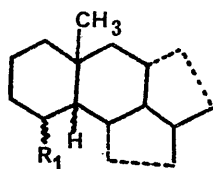
The germacranolides are particularly important because they are believed to be intermediates in the biogenesis of some other sesquiterpenes (e.g. guianolides). The x-ray analyses of the two sesquiterpenes in this chapter have been carried out to provide conformational information both to assist in biogenetic investigations and as part of a study of medium ring dienes.

2.2 Costunolide

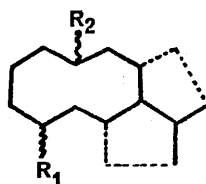
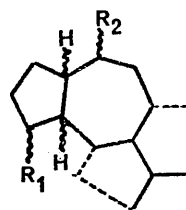
Costunolide (I), a germacranolide, has been extracted from the plant, Saussurea lappa Clerke (4), found in the Himalayas and from Artemisia balchanorum H. Krasch (5), which occurs in Central Asia. Professor Kupchan, (University of Virginia) kindly supplied the sample of costunolide used in the analysis which follows.

Previous work by diffraction (5, 6), synthetic (7) and IR (8) methods on costunolide and its silver nitrate adduct failed to yield the complete structure. The evidence from these studies for the proposed conformation (Fig. 2.2) was supported by the work of Tori et al (9)

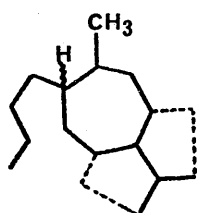
Anolide



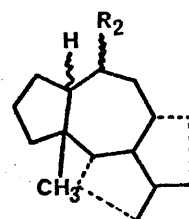
Guianolide



Germacranolide



Xantholide

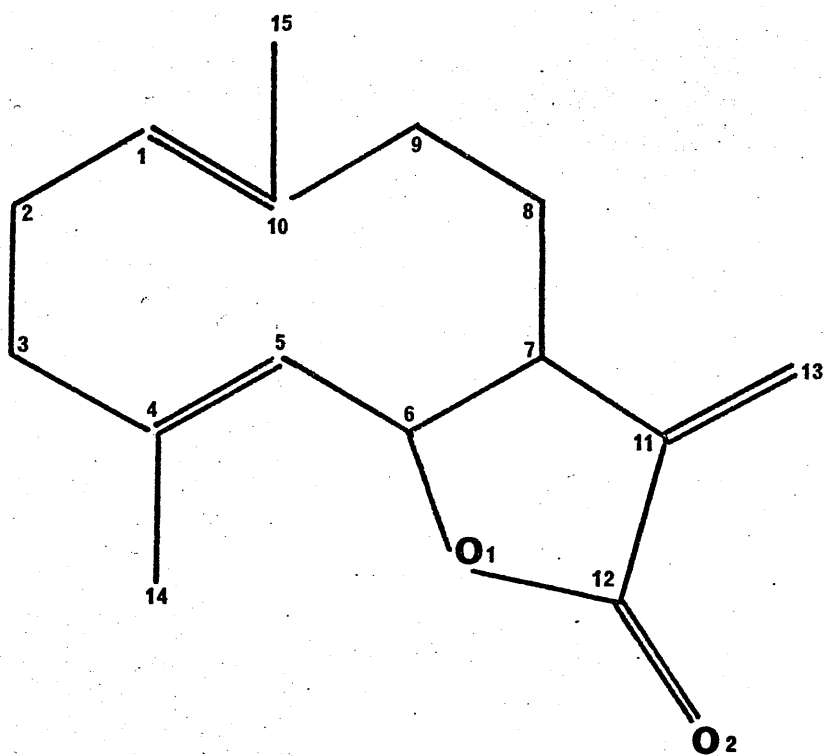


Ambrosanolide

R₁ = CH₃ or H

R₂ = CH₃ or H

Fig.2.1 Sesquiterpenoids: Most frequently occurring skeletons



(I)

Fig. 2.2 Structure of Costunolide

based on the simultaneous application of the nuclear overhauser effect and an NMR shift reagent, and by the ORD/CD results of Suchy and co-workers (10).

The earlier x-ray analyses (5,6) had used Mo-K α radiation for four circle diffractometer measurement of intensities for costunolide (6) and its silver nitrate adduct (5) but had not revealed the hydrogen atoms. Further, the effect of the silver nitrate on the conformation of the macrocycle was not clear. It was decided to recollect intensity data using Cu-K α radiation in the hope that some of the weaker reflexions could be included and thus provide a more accurate structure.

2.3 Experimental

Crystal data

Costunolide $C_{15}H_{20}O_2$

Unit cell dimensions $a = 11.028 \text{ \AA}$
 $b = 15.400 \text{ \AA}$ $\alpha = \beta = \gamma = 90^\circ$
 $c = 7.987 \text{ \AA}$

Space group $P2_12_12_1 (D_2^4)$

$M = 232 \text{ a.m.u.}$

$V = 1356 \text{ \AA}^3$

$D_o = 1.159 \text{ gm cm}^{-3}$

$D_c = 1.136 \text{ gm cm}^{-3}$

$\mu = 5.9 \text{ cm}^{-1} (\text{Cu-K}\alpha)$

$F(000) = 504$

$Z = 4$

The space group was determined uniquely by the systematic absences:

h00 when $h=2n+1$, 0k0 when $k=2n+1$, 00l when $l=2n+1$.

The crystal used in this analysis was a small cube with a 1mm side.

Data Collection

Cell measurements were available from the previous analysis and these were confirmed from precession photographs. The crystal was transferred to a Hilger & Watts Y290 four circle diffractometer controlled by a PDP-8 computer. Cu-K α radiation filtered through nickel was employed.

The positions of the reciprocal axes with respect to the crystal faces were approximately known from the photographic work and a starting pair of reflexions was obtained by scanning the Φ circle at 2θ values calculated from the cell dimensions. These reflexions were optimised by 2θ , Φ and χ scans and indexed to obtain an initial orientation matrix. This orientation matrix was then used to calculate the angular settings for a further 15 reflexions which were also optimised as described above. The unit cell parameters were refined using these 17 setting angles in a least squares procedure and are quoted in the crystal data.

Intensity measurements were made with the θ , 2θ scan procedure. Typical scan parameters were as follows:

Increment in θ = 0.01°

No. of scan steps = 80

Time per step = 1 sec.

Background counts were made for 20 seconds at each extreme position of the scan.

The integrated intensity, \underline{I} , was calculated from the peak, \underline{P} , and background $\underline{B1}$, $\underline{B2}$, counts using equation (1):

$$I = P - (B1 + B2) \cdot \frac{t_p}{2t_b} \quad (1)$$

t_p = counting time at peak.

t_b = " " for one of the background counts (both the same).

and the standard deviation of \underline{I} , (σI), was obtained from equation (2):

$$\sigma I = \left(P + (B1 + B2) \cdot \frac{t_p}{2t_b} \right)^{\frac{1}{2}} \quad (2)$$

Two strong reflexions were used as standards throughout the data collection. Their counts were checked after every 20 intensity measurements. These monitor counts held constant to within about 2 % during the data collection and were used to place all the measurements on a common scale.

Reflexion intensities were collected for $2\theta \leq 110^\circ$ in the two equivalent octants $\underline{hkl}, \underline{h\bar{k}\bar{l}}$. Data for which $I \leq 2\sigma I$ were discarded and equivalent reflexions were averaged to give 1398 independent structure amplitudes.

A number of reflexions had high counts which may have saturated the counter. Accordingly data for which $2\theta \leq 20^\circ$ were recollected with a reduced beam intensity so that a counting loss correction could be applied.

Structure Analysis

A trial set of non-hydrogen co-ordinates was available from the Mo-K α analysis and these gave an initial residual R of 25.2% with fixed isotropic temperature factors. Four cycles of least squares refinement of these co-ordinates, allowing for anisotropic thermal motion, reduced R to 14.4%. A difference synthesis was carried out and this yielded most of the hydrogen atoms. These hydrogen locations were refined with invariant isotropic temperature factors during a further two cycles of least squares calculations. A new difference synthesis yielded the remaining hydrogen atoms. Two further cycles of full matrix least squares adjustment, during which the isotropic thermal parameters of the hydrogen atoms were also refined, converged at $R = 7.9\%$

Counting loss errors were then corrected and this gave an R of 6.8%. Finally a weighting scheme of the form:

$W = x.y$ Where the weight, W , was calculated
as follows,

(a) For $0.7|F_o| > |F_c|$, $W = 1 \times 10^{-9}$ otherwise,

(b) $x = \sin\theta/2.1$

(c) $y = 1$, unless $|F_o| > 3.05$ when $y = 3.05/|F_o|$

(x and y were selected to minimize the deviation from constant $W\Delta^2$ over the whole range of $\sin\theta$ and $|F_o|$).

was applied and after two cycles of least squares calculations a value of $R=4.3\%$ resulted. The ratio of observations to parameters was 6.0 to 1 and no correction was made for absorption.

2.4 Results

Table I Costunolide: Fractional atomic coordinates

Positional estimated standard deviations are shown in parentheses.

The hydrogen atoms are numbered according to the atoms to which they are attached.

The table shows:

Atom	x	y	z,
------	---	---	----

C(1)	0876(2)	6935(1)	1874(3)
C(2)	-0116(2)	6850(2)	3117(4)
C(3)	0073(2)	6014(2)	4177(4)
C(4)	1345(2)	6021(1)	4881(3)
C(5)	2222(2)	5671(1)	3959(2)
C(6)	3533(2)	5880(1)	4067(2)
C(7)	4125(2)	6064(1)	2351(2)
C(8)	4083(2)	7009(1)	1742(3)
C(9)	2919(2)	7313(1)	0836(3)
C(10)	1841(2)	7448(1)	1951(2)
C(11)	5381(2)	5709(1)	2628(3)
C(12)	5322(2)	5093(1)	4052(4)
C(13)	6414(3)	5851(2)	1855(5)
C(14)	1566(3)	6526(2)	6435(3)
C(15)	1967(3)	8182(2)	3166(4)
O(1)	4218(1)	5144(1)	4768(2)
O(2)	6080(2)	4591(1)	4551(4)
H(1)	0951(22)	6544(16)	1014(32)
H(21)	-0049(26)	7476(19)	3876(37)
H(22)	-0944(29)	6786(22)	2608(45)
H(31)	-0574(32)	5978(23)	5088(55)
H(32)	-0114(39)	5484(27)	3342(58)
H(5)	1972(21)	5289(14)	2937(30)
H(6)	3712(20)	6360(15)	4868(32)
H(7)	3717(26)	5699(20)	1526(39)
H(81)	4256(30)	7465(23)	2759(44)
H(82)	4772(29)	7079(21)	1043(41)
H(91)	3098(24)	7905(17)	0222(35)
H(92)	2775(22)	6848(16)	-0001(33)
H(131)	7251(39)	5485(27)	2284(59)
H(132)	6429(19)	6113(14)	1318(29)
H(141)	2401(29)	6544(20)	6770(42)
H(142)	1045(30)	6359(23)	7281(44)
H(143)	1362(32)	7123(25)	6224(51)
H(151)	1236(43)	8261(32)	3903(64)
H(152)	2435(38)	8662(30)	2776(58)
H(153)	2492(53)	8000(38)	4211(73)

Table II Costunolide: Thermal parameters

Carbon and oxygen values are anisotropic.

Hydrogen values are isotropic.

Estimated errors in the last two digits are shown in parentheses.

All figures are multiplied by 10^4 .

The anisotropic temperature factor expression used was of the form described by equation (12) in chapter one, the coefficients being in terms of the mean square amplitudes of vibration U_{ij} .

The table shows:

Atom	U_{11} or U	U_{22}	U_{33}	U_{12}	U_{13}	U_{23}
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C(1)	824(12)	800(11)	734(11)	21(10)	-172(10)	60(10)
C(2)	693(11)	1059(16)	1112(18)	81(11)	-80(13)	204(15)
C(3)	676(11)	1014(16)	1148(18)	51(11)	147(12)	253(15)
C(4)	735(10)	770(10)	751(11)	77(9)	141(9)	201(10)
C(5)	683(9)	615(9)	698(10)	-6(8)	94(9)	64(8)
C(6)	684(9)	579(8)	618(9)	29(7)	26(8)	34(7)
C(7)	697(9)	645(9)	618(9)	-85(8)	39(8)	0(8)
C(8)	798(12)	796(12)	883(14)	-144(10)	23(11)	213(12)
C(9)	1011(14)	878(13)	694(11)	-42(12)	7(12)	248(10)
C(10)	898(12)	676(9)	645(9)	-14(10)	-123(9)	107(9)
C(11)	704(10)	695(10)	883(13)	-41(8)	120(10)	-97(10)
C(12)	698(10)	763(11)	1095(16)	64(10)	-6(12)	20(12)
C(13)	894(16)	1169(20)	1444(25)	-88(15)	358(18)	110(21)
C(14)	1059(17)	1209(19)	702(13)	289(16)	208(12)	45(13)
C(15)	1381(23)	818(14)	1061(18)	-198(15)	48(19)	-117(14)
O(1)	773(7)	809(8)	889(9)	137(6)	66(7)	236(8)
O(2)	875(10)	1078(12)	1754(22)	289(10)	-13(13)	301(15)
H(1)	590(61)					
H(21)	822(81)					
H(22)	897(96)					
H(31)	884(114)					
H(32)	1074(143)					
H(5)	458(57)					
H(6)	440(57)					
H(7)	584(79)					
H(81)	937(98)					
H(82)	827(87)					
H(91)	694(68)					
H(92)	624(64)					
H(131)	1287(146)					
H(132)	563(53)					
H(141)	781(89)					
H(142)	859(100)					
H(143)	935(112)					
H(151)	1267(159)					
H(152)	1238(132)					
H(153)	1670(210)					

TABLE III

COSIUNOLIDE: INTRAMOLECULAR BONDED DISTANCES (ANGSTROMS)
STANDARD DEVIATIONS (ANGSTROMS) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	DIST.	ATOM1	ATOM2	DIST.
C1	C2	1.484 (3)	C2	H21	1.123 (27)
C1	C10	1.326 (2)	C2	H22	1.008 (32)
C2	C3	1.555 (3)	C3	H31	1.033 (39)
C3	C4	1.512 (3)	C3	H32	1.052 (37)
C4	C5	1.329 (2)	C5	H5	1.046 (24)
C4	C14	1.486 (3)	C6	H6	0.988 (21)
C5	C6	1.483 (2)	C7	H7	0.972 (27)
C6	C7	1.544 (3)	C8	H81	1.081 (38)
C6	O1	1.473 (1)	C8	H82	0.942 (33)
C7	C8	1.536 (2)	C9	H91	1.056 (23)
C7	C11	1.506 (2)	C9	H92	0.997 (26)
C8	C9	1.546 (3)	C13	H131	1.113 (38)
C9	C10	1.500 (3)	C13	H132	0.665 (27)
C10	C15	1.497 (3)	C14	H141	0.976 (28)
C11	C12	1.482 (3)	C14	H142	0.931 (36)
C11	C13	1.313 (3)	C14	H143	0.952 (25)
C12	O1	1.348 (2)	C15	H151	1.006 (47)
C12	O2	1.205 (2)	C15	H152	0.989 (37)
C1	H1	0.978 (29)	C15	H153	1.035 (64)

TABLE IV COSYNOCLIDE: INTERMOLECULAR CONTACTS OF LESS THAN 4.0 ANGSTROMS

ATOM1	ATOM2	DIST #	ATOM1	ATOM2	DIST #
C(3)....C(4)	*	3.859	O(2)....C(2)	*	3.764
C(3)....C(13)	*	3.938	O(2)....C(13)	**	3.389
C(4)....C(7)	*	3.803	C(8)....C(14)	**	3.835
C(4)....C(11)	*	3.941	C(12)....C(15)	**	3.910
C(14)....C(12)	*	3.864	C(9)....O(2)	****	3.691
C(14)....O(1)	*	3.802	C(15)....O(2)	****	3.748
O(1)....C(1)	*	3.618	C(8)....C(1)	*****	3.860
O(1)....C(5)	*	3.912	C(13)....C(9)	*****	3.921
O(2)....C(1)	*	3.691			

IN ANGSTROMS

THE ASTERISKS(*) REFER TO THE FOLLOWING TRANSFORMATIONS OF THE ATOMIC COORDINATES

* 0.5-X, 1.0-Y, 0.5+Z
 ** 1.5-X, 1.0-Y, 0.5+Z
 *** 0.5+X, 1.5-Y, 1.0-Z
 **** 1.0-X, 0.5+Y, 0.5-Z
 ***** 0.5+X, 1.5-Y, -Z

TABLE V

COSIUNOLIDE: VALENCY ANGLES(DEGREES)
STANDARD DEVIATIONS(DEGREES) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	ATOM3	ANGLE	ATOM1	ATOM2	ATOM3	ANGLE
C10	C1	C2	127.7(2)	C3	C2	C1	109.7(1)
C4	C3	C2	108.8(1)	C5	C4	C3	117.8(2)
C14	C4	C3	117.8(2)	C14	C4	C5	123.8(1)
C6	C5	C4	126.1(1)	C7	C6	C5	113.7(1)
O1	C6	C5	110.8(1)	O1	C6	C7	105.2(1)
C8	C7	C6	116.2(1)	C11	C7	C6	101.1(1)
C11	C7	C8	114.8(1)	C9	C8	C7	117.4(1)
C10	C9	C8	115.0(2)	C15	C10	C1	123.8(2)
C9	C10	C1	121.7(1)	C15	C10	C9	114.5(1)
C12	C11	C7	107.8(1)	C13	C11	C7	131.8(2)
C13	C11	C12	120.4(2)	O1	C12	C11	109.1(1)
O2	C12	C11	129.2(2)	O2	C12	O1	121.7(2)
C12	O1	C6	110.4(1)	H1	C1	C2	119.9(14)
H1	C1	C10	111.5(14)	H22	C2	C1	114.7(24)
H21	C2	C1	103.2(16)	H21	C2	C3	113.7(18)
H22	C2	C3	106.0(16)	H22	C2	H21	109.8(21)
H32	C3	C2	104.4(21)	H31	C3	C4	112.6(22)
H32	C3	C4	113.8(18)	H31	C3	C2	108.8(14)
H32	C3	H31	108.1(26)	H5	C5	C4	118.1(11)
H5	C5	C6	114.9(11)	H6	C6	O1	103.4(12)
H6	C6	C5	112.5(11)	H6	C6	C7	110.6(13)
H7	C7	C11	108.5(12)	H7	C7	C6	106.7(16)
H7	C7	C8	109.0(14)	H81	C8	C9	108.4(16)
H82	C8	C9	110.6(21)	H82	C8	C7	105.1(13)
H81	C8	C7	110.5(18)	H82	C8	H81	103.9(26)

VALENCE ANGLES (CONTINUED)

ATOM1	ATOM2	ATOM3	ANGLE	ATOM1	ATOM2	ATOM3	ANGLE
H92	C9	C10	112.1(14)	H91	C9	C8	109.0(14)
H92	C9	C8	103.0(13)	H91	C9	C10	107.0(15)
H92	C9	H91	110.7(24)	H131	C13	C11	119.3(25)
H132	C13	C11	119.9(19)	H132	C13	H131	120.8(31)
H142	C14	C4	111.5(20)	H143	C14	C4	107.3(28)
H141	C14	C4	113.2(20)	H142	C14	H141	113.0(31)
H143	C14	H141	104.7(22)	H143	C14	H142	106.5(28)
H152	C15	C10	114.7(30)	H153	C15	C10	112.7(24)
H151	C15	C10	111.6(23)	H152	C15	H151	124.0(28)
H153	C15	H151	91.0(44)	H153	C15	H152	98.5(39)

TABLE VI

COSIUMOLIDE: TORSION ANGLES (DEGREES)
STANDARD DEVIATIONS (DEGREES) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	ATOM3	ATOM4	ANGLE	ATOM1	ATOM2	ATOM3	ATOM4	ANGLE
C10	C1	C2	C3	-101.9(2)	C2	C1	C10	C9	164.7(2)
C2	C1	C10	C15	-13.5(3)	C1	C2	C3	C4	52.2(2)
C2	C3	C4	C5	-88.7(2)	C2	C3	C4	C14	82.8(2)
C3	C4	C5	C6	155.9(1)	C14	C4	C5	C6	-15.0(3)
C4	C5	C6	C7	-131.2(1)	C4	C5	C6	O1	110.6(2)
C5	C6	C7	C8	88.7(1)	C5	C6	C7	C11	-146.4(1)
O1	C6	C7	C8	-149.9(1)	O1	C6	C7	C11	-25.0(1)
C5	C6	O1	C12	145.1(1)	C7	C6	O1	C12	21.9(1)
C6	C7	C8	C9	-84.0(2)	C11	C7	C8	C9	158.4(1)
C6	C7	C11	C12	20.5(1)	C6	C7	C11	C13	-159.8(2)
C8	C7	C11	C12	146.4(1)	C8	C7	C11	C13	-33.9(3)
C7	C8	C9	C10	73.6(2)	C8	C9	C10	C1	-111.7(2)
C8	C9	C10	C15	66.6(2)	C7	C11	C12	O1	-8.5(2)
C7	C11	C12	O2	170.0(2)	C13	C11	C12	O1	171.8(2)
C13	C11	C12	O2	-9.7(4)	C11	C12	O1	C6	-8.7(2)
O2	C12	O1	C6	172.7(2)	C10	C1	C2	H21	19.6(15)
C10	C1	C2	H22	139.0(20)	H1	C1	C2	C3	66.4(17)
H1	C1	C2	H21	-172.1(23)	H1	C1	C2	H22	-52.7(27)
H1	C1	C10	C9	-4.4(16)	H1	C1	C10	C15	177.4(16)
C1	C2	C3	H31	175.2(19)	C1	C2	C3	H32	-69.6(20)
H21	C2	C3	C4	-62.8(16)	H21	C2	C3	H31	60.2(25)
H21	C2	C3	H32	175.4(26)	H22	C2	C3	C4	176.5(19)
H22	C2	C3	H31	-60.5(27)	H22	C2	C3	H32	54.7(28)
H31	C3	C4	C5	150.6(19)	H31	C3	C4	C14	-37.9(19)
H32.	C3	C4	C5	27.1(21)	H32	C3	C4	C14	-161.3(21)

TORSION_ANGLES (CONTINUED)

ATOM1	ATOM2	ATOM3	ATOM4	ANGLE	ATOM1	ATOM2	ATOM3	ATOM4	ANGLE
C3	C4	C5	H5	-12.4(13)	C14	C4	C5	H5	176.6(13)
C3	C4	C14	H141	-176.0(18)	C3	C4	C14	H142	55.3(20)
C3	C4	C14	H143	-61.0(21)	C5	C4	C14	H141	-5.0(19)
C5	C4	C14	H142	-133.7(20)	C5	C4	C14	H143	110.0(21)
C4	C5	C6	H6	-4.6(13)	H5	C5	C6	C7	37.4(12)
H5	C5	C6	O1	-80.8(12)	H5	C5	C6	H6	164.1(18)
C5	C6	C7	H7	-33.1(15)	O1	C6	C7	H7	88.3(15)
H6	C6	C7	C8	-39.0(12)	H6	C6	C7	C11	86.0(12)
H6	C6	C7	H7	-160.7(19)	H6	C6	O1	C12	-94.2(12)
C6	C7	C8	H81	41.0(19)	C6	C7	C8	H82	152.6(19)
C11	C7	C8	H81	-76.6(19)	C11	C7	C8	H82	35.0(19)
H7	C7	C8	C9	36.5(15)	H7	C7	C8	H81	161.5(24)
H7	C7	C8	H82	-86.9(24)	H7	C7	C11	C12	-91.4(15)
H7	C7	C11	C13	88.2(15)	C7	C8	C9	H91	-166.4(15)
C7	C8	C9	H92	-48.7(14)	H81	C8	C9	C10	-52.5(19)
H81	C8	C9	H91	67.5(24)	H81	C8	C9	H92	-174.8(24)
H82	C8	C9	C10	-165.8(19)	H82	C8	C9	H91	-45.8(25)
H82	C8	C9	H92	71.9(24)	H91	C9	C10	C1	127.2(15)
H91	C9	C10	C15	-54.5(15)	H92	C9	C10	C1	5.6(15)
H92	C9	C10	C15	-176.1(15)	C1	C10	C15	H151	-3.0(26)
C1	C10	C15	H152	-150.7(27)	C1	C10	C15	H153	97.7(32)
C9	C10	C15	H151	178.7(26)	C9	C10	C15	H152	31.1(27)
C9	C10	C15	H153	-80.5(32)	C7	C11	C13	H131	-177.6(24)
C7	C11	C13	H132	3.2(24)	C12	C11	C13	H131	2.1(25)

THE ANGLE 1-2-3-4 IS DEFINED AS POSITIVE IF WHEN VIEWED ALONG THE 2-3 BOND, ATOM 1 HAS TO BE ROTATED CLOCKWISE TO ECLIPSE ATOM 4

TABLE VII COSTUNOLIDE: MEAN PLANE CALCULATIONS

	ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
1.	C(1) C(2) C(10)	C(15) H(1)	0.000 (2) 0.000 (3) 0.000 (2) -0.291 (3) 0.156 (25)
2.	C(1) C(9) C(10)	C(15) H(1)	0.000 (2) 0.000 (2) 0.000 (2) 0.043 (3) -0.076 (25)
3.	C(3) C(4) C(5)	C(14) H(5)	0.000 (3) 0.000 (2) 0.000 (2) -0.197 (3) 0.215 (23)
4.	C(4) C(5) C(6)	C(14) H(5)	0.000 (2) 0.000 (2) 0.000 (2) 0.317 (3) -0.168 (23)

TABLE VII (CONTINUED)

	ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
5.	C(11) C(12) O(1)	C(6) C(7) C(13) O(2)	0.000(2) 0.000(3) 0.000(2) 0.212(2) -0.209(2) 0.158(4) -0.028(2)
6.	C(7) C(11) C(12)	C(6) C(13) O(1) O(2)	0.000(2) 0.000(2) 0.000(3) 0.531(2) -0.008(4) 0.186(2) -0.164(2)
7.	C(13) H(131) H(132)	C(11)	0.000(4) 0.000(44) 0.000(22) 0.055(2)

TABLE VII (CONTINUED)

ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
C(11)		-0.003(2)
C(12)		0.008(3)
O(1)		-0.003(2)
O(2)		-0.003(2)
	C(6)	0.191(2)
	C(7)	-0.231(2)
	C(13)	0.163(4)

THE OUT OF PLANE ATOMS WERE NOT INCLUDED IN THE DERIVATION OF THE PLANES.

THE EQUATIONS OF THESE PLANES ARE:

1. $0.0510X - 0.7222Y + 0.4672Z = -6.5200$
2. $0.4081X - 0.6021Y + 0.6863Z = -5.0081$
3. $-0.1662X - 0.8870Y + 0.4308Z = -6.7909$
4. $0.1196X - 0.7199Y + 0.6837Z = -3.8317$
5. $0.3347X + 0.7143Y + 0.6146Z = 9.5561$
6. $0.1939X + 0.7481Y + 0.6347Z = 9.0594$
7. $0.1829X + 0.7115Y + 0.6784Z = 8.7100$
8. $0.3448X + 0.7064Y + 0.6182Z = 9.5564$

2.5 Discussion of Results

The results of the analysis are shown in Tables I - VII. Figs. 2.3 and 2.4 illustrate a view of the molecule, and of the crystal packing viewed along the c axis, respectively. The final and observed structure factors are listed in Table XIIa.

The extended chair conformation of the cyclodecadiene ring is very similar to that in a number of related sesquiterpenes (Fig. 2.5) as indicated in Table VIII. Strain energy calculations (15) have shown this to be the minimum energy conformation of trans, trans - cyclodeca-1,5-diene. The methyl substituents C(14) and C(15) are cis and β. This arrangement has also been observed in other germacranes e.g. eupatolide (11), alatolide (12), eupatoriopicrin (13), onopordopicrine (14), pregeijerene (16), germacratrine (17) and elephantol (18). Cis and α-(19) and anti- (20) stereochemistries are also known.

The C- C(sp³) - C valency angles in the decadiene ring vary from 109° to 117° with a mean of 113.5°. These large deviations from the tetrahedral angle are indicative of considerable strain. This situation is quite common in saturated medium ring compounds many of which exhibit valency angles of 116 - 117° (21), and angles in excess of the tetrahedral value are not unusual (Table IX).

There is distortion of the double bonds at C(1) - C(10) and C(4) - C(5) as the respective torsion angles of 164° and 156° exhibit a marked departure from planarity. Newman projections down these two bonds are shown in Fig. 2.6 and Table X describes the deformations in costumlide and some related structures in terms of τ (torsion) - and

Fig. 2.3 Costunolide: A view of the molecule



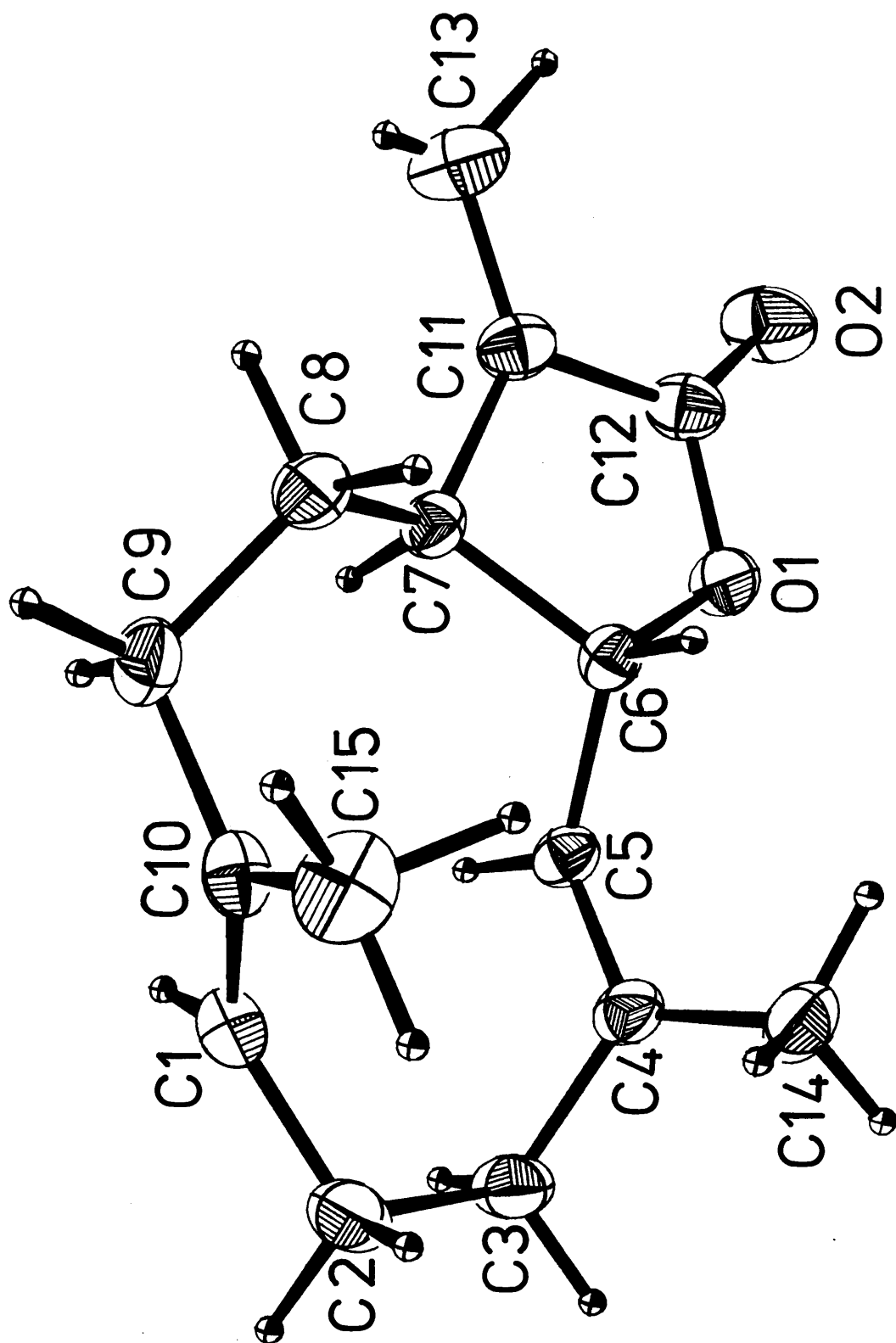
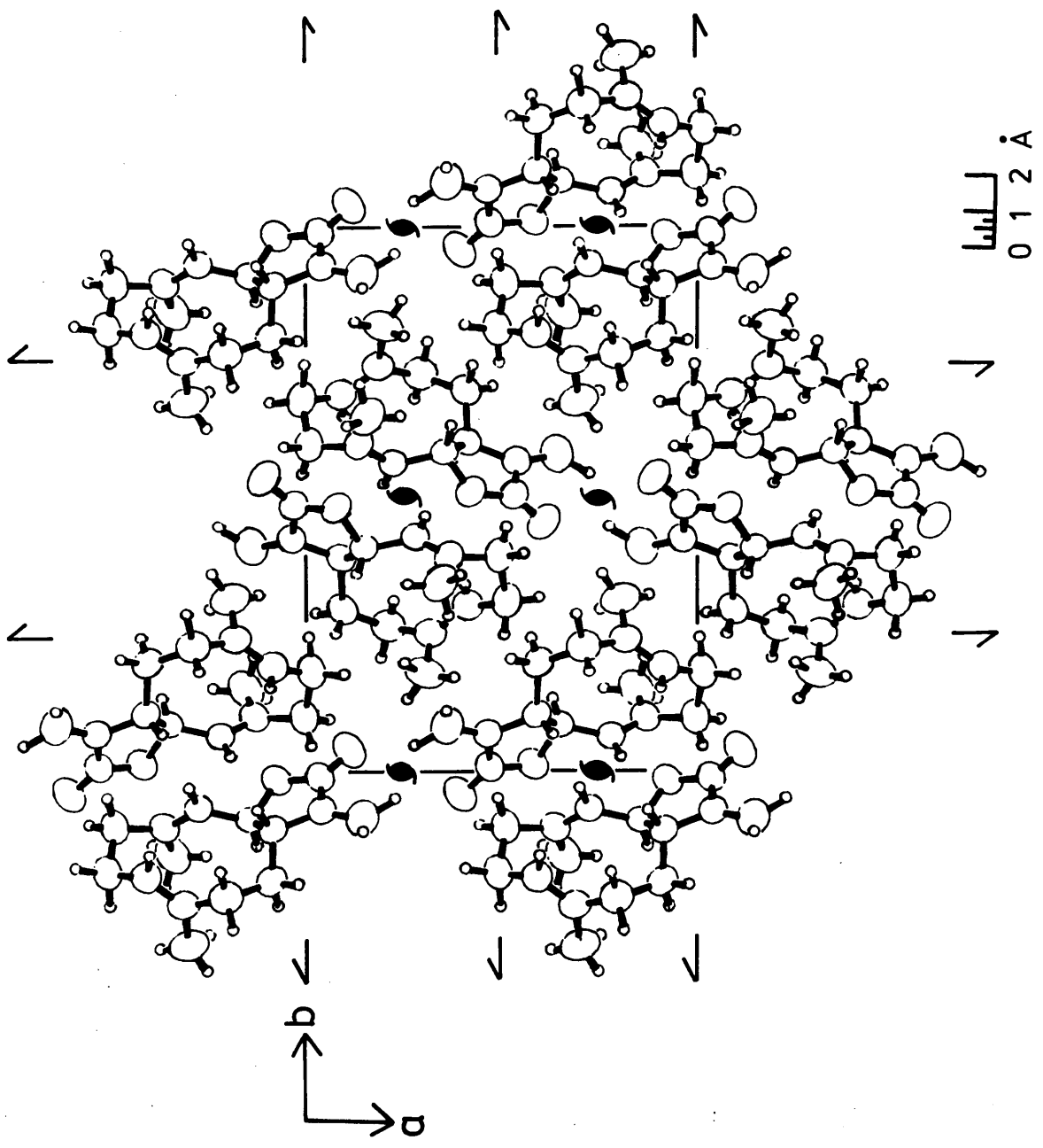
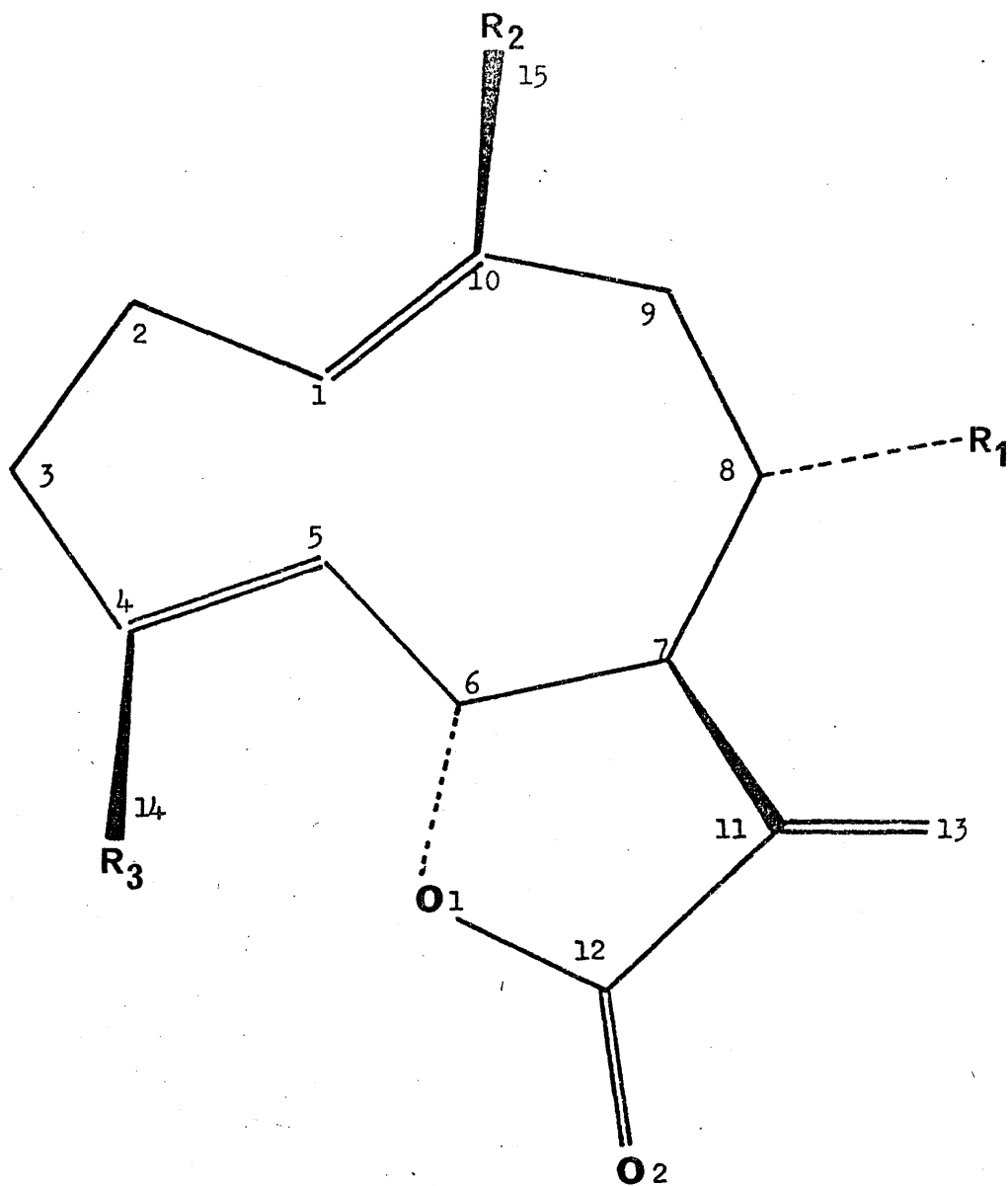


Fig. 2.4 Costunolide crystal packing: A view down the 'c' axis





	R_1	R_2	R_3
Costunolide	H	CH_3	CH_3
Eupatolide	OH	CH_3	
Alatolide	$OCOCH(CH_3)_2$	CH_2OH	CH_2OH
Eupatoriopicrin	$OCOC(CH_2OH)CHCH_2OH$	CH_3	CH_3
Onopordopicrine	$OCOC(CH_2)CH_2OH$	CH_3	CH_2OH
Jurineolide *	$OCOC(CHCH_2OH)CH_3$	CH_2OH	CH_2OH
Albicolide *	H	CH_2OH	CH_2OH

* No x-ray analysis currently available. Stereochemistry based on ORD/CD observations.

Fig. 2.5 Structure of Costunolide and some related Sesquiterpenoids

TABLE VIII MACROCYCLE TORSION ANGLES IN VARIOUS GERMACRANOLIDES

		COMPOUND					
TORSION ANGLE		CO	EL	AL	ER	ON	FF
1 - 2		-102	-98	-107	-102	-100	-96
2 - 3		52	50	50	50	53	44
3 - 4		-89	-86	-83	-87	-82	-82
4 - 5		156	155	154	154	153	168
5 - 6		-131	-136	-137	-133	-141	-132
6 - 7		89	90	93	90	95	77
7 - 8		-84	-76	-87	-77	-83	-87
8 - 9		74	67	76	66	70	78
9 - 10		-112	-115	-108	-109	-115	-113
10 - 1		164	167	167	166	168	167
CO	COSTUNOLIDE						
AL	ALATOLIDE (12)			EL		EUPATOLIDE (11)	
ON	ONOPORDOPICRINE (14)			ER		EUPATORIOPICRIN (13)	
				FF		CALC. DIMETHYL CYCLODECADIENE	

TORSION ANGLES (DEGREES). NUMBERING AS IN FIG. 2.5

TABLE IX MACROCYCLE C-C(SP³3)-C VALENCE ANGLES IN GERMACRANOLIDES

	RANGE		MEAN
COSTUNOLIDE	109	TO 117	113.5
EUPATOLIDE (11)	109	TO 118	114.8
ALATOLIDE (12)	109	TO 115	112.7
EUPATORIOPICRIN (13)	108	TO 119	114.8
ONOPORDOPICRINE (14)	107	TO 129	119.0
PREGEIJERENE (16) *	106	TO 125	118.1
GERMACRATRIENE (17) *	102	TO 116	109.2
ELEPHANTOL (18)	107	TO 124	113.9
DIHYDROMIKANOLIDE (19)	115	TO 122	117.7

ANGLES IN DEGREES

* A GERMACRANE DERIVATIVE CONTAINING NO LACTONE FUNCTION

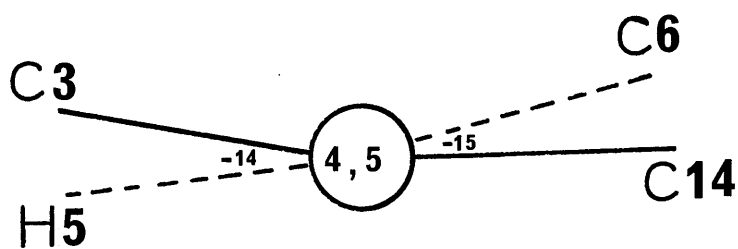
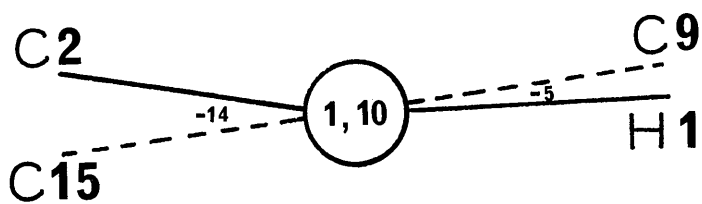


Fig. 2.6 Newman projections down the C(1)-C(10) and C(4)-C(5) bonds of Costunolide

TABLE X THE DISTORTION OF THE CYCLODECAADIENE DOUBLE BONDS IN SOME
GERMACRANOLIDES

	C(3) - C(4) - C(5) - C(6)			C(9) - C(10) - C(1) - C(2)				
	PHI1	CHI1	CHI2	TOR1	PHI2	CHI3	CHI4	TOR2
COSTUNOLIDE	156.0	-9.1	10.5	165.8	164.5	-1.9	10.5	170.6
EUPATOLIDE(11)	154.9	-8.9	17.7	168.2	166.6	-1.3	14.5	174.5
ALATOLIDE(12)	153.5	-6.1	5.5	174.2	166.6	0.4	25.2	179.0
ONOPORDOPICRINE(14)	153.4	-10.0	12.5	169.7	168.0	-1.9	12.9	175.5
CALCULATED *	168.3	-5.8	5.9	173.5	166.7	-4.3	4.3	173.2

* FOR DIMETHYL CYCLODECAADIENE

PHI1	TORSION ANGLE C(3)-C(4)-C(5)-C(6)
PHI2	TORSION ANGLE C(9)-C(10)-C(1)-C(2)

THE FOLLOWING SYMBOLS DESCRIBE THE DISTORTION PARAMETERS OF WINKLER AND
DUNITZ(23)

CHI1	OUT OF PLANE BENDING OF C(14)
CHI2	OUT OF PLANE BENDING OF H(5)
CHI3	OUT OF PLANE BENDING OF C(15)
CHI4	OUT OF PLANE BENDING OF H(1)
TOR1	TWIST ABOUT C(4)-C(5)
TOR2	TWIST ABOUT C(10)-C(1)

χ (out of plane) - bending according to the method of Winkler and Dunitz (23). The relative uncertainty of the locations of the hydrogens at C(1) and C(5) prohibits detailed discussion of the values in Table X. It is apparent however that τ and χ are relatively constant from molecule to molecule with χ for the unsubstituted trigonal atoms exhibiting the widest range of values. The deformations at C(4) - C(5) result from almost equal amounts of torsion and out of plane bending while at the other double bond there is virtually no out of plane bending at C(10). The transannular separations C(1)....C(4) and C(1)....C(5) are much shorter ($2.8 - 2.9\text{\AA}$) than C(10)....C(4) and C(10)...C(5) ($3.2 - 3.4\text{\AA}$) and the van der Waals repulsion between C(1), C(4), C(5) and between their substituents is probably partially relieved by the out of plane bending of H(1), C(14) and H(5) which is in a direction outwards from the ring. The absence of a similar deformation at C(10) may be related to the less crowded environment in the vicinity of this atom.

The C(1) - C(5) trans-annular distance of 2.96\AA is very short and this is not unexpected when the separations found for some other germacrane are compared (Table XI). Costunolide shows a UV absorption at $220\text{nm}(+ve)$ (10) and other compounds e.g. jurineolide (10), albicolide (10), have a similar UV band at about 220 nm. It is believed (10, 24) that such absorption at about 200 and 220 nm is the result of a trans-annular interaction between the double bonds of the macrocycle giving rise to a $\Pi \rightarrow \Pi^*$ transition and that the sign of the absorption reflects the chirality of the diene. The spectra of costunolide, jurineolide, albicolide, and alatolide are all very

TABLE XI TRANSANNULAR SEPARATION IN SOME GERMACRANOLIDES

	DIST +
CUSTUNOLIDE	2.96
EUPATOLIDE (11)	2.94
ALATOLIDE (12)	2.88
EUPATORIOPICRIN (13)	2.94
ONOPORDOPICRINE (14)	2.89
PREGEIJERENE (16)	2.91
ELEPHANTOL (18)	2.98
DIHYDROMIKANOLIDE (19)	3.03
CALCULATED *	2.84

C(1)...C(5) TRANSANNULAR SEPARATION IN ANGSTROMS

FOR DIMETHYL CYCLODECADIENE

+

*

similar in this region and this implies that they have identical stereochemistry in the vicinity of the double bonds. The short trans-annular distances, the observed chiral relationship of the double bonds, and the UV spectra all support the case for the postulated intermediate role of germacrane derivatives in biogenesis (1).

The other major feature of the CD spectrum of costunolide (Fig. 2.7) is the negative Cotton effect at 261 nm (7). This has been attributed to the $n \rightarrow \pi^*$ transition of the $C=C-C=O$ system (25, 26). Stöcklin et al (3) have correlated the sign of the absorption with the position and stereochemistry of the lactone ring fusion. In the case of a 6,7 trans-fusion (Fig. 2.3) a negative Cotton effect is predicted in complete agreement with the observed spectrum. Beecham (27) has proposed that there is a relationship between this sign and the $C=C-C=O$ torsion angle $\underline{\psi}$. The value of this angle for costunolide is -10° which, according to Beecham, would give a negative sign. The view has recently been put forward (12) that the $C_{\alpha}-C_{\beta}-C_{\gamma}-O$ torsion angle $\underline{\phi}$ of the $\underline{\alpha}$ -methylene- $\underline{\gamma}$ -lactone may have more relevance. This suggestion is based on the facts that (a) the observed variation in $\underline{\psi}$ is large with little apparent effect on the CD spectrum, (b) $\underline{\psi}$ and $\underline{\phi}$ are nearly always in the same sense with the latter usually having greater magnitude, and (c) the more reliable predictive powers of Stöcklin's rule when contrasted with (a)

$\Delta\epsilon$

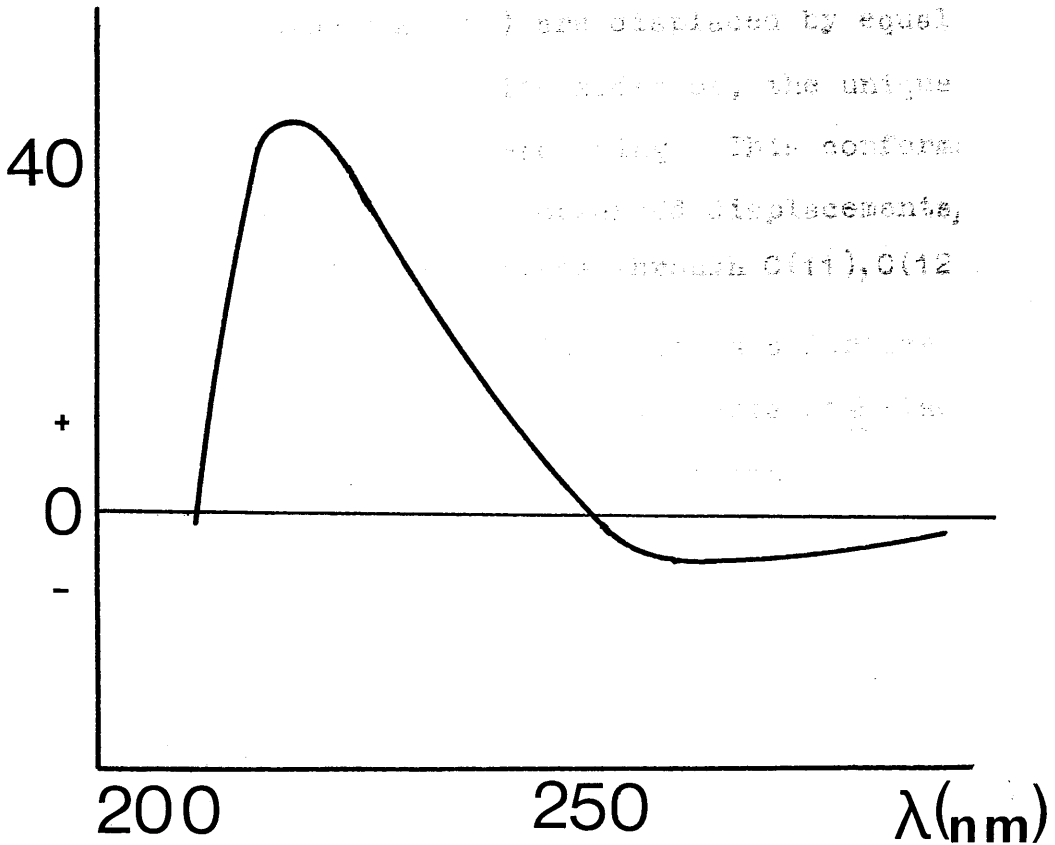


Fig. 2.7 CD spectrum of Costunolide

A quantitative description of the half-chair conformation of the δ -lactone ring was obtained by the method of Cremer and Pople (28). This gave $\underline{q} = 0.25A^\circ$, $\underline{\Phi} = 17^\circ$. Since the alternate half-chairs and envelopes assumed by an isolated five-membered ring are separated by phase angles of 18° the conformation adopted by the fused δ -lactone in costunolide is shown as being almost exactly the half-chair in which C(6) and C(7) are displaced by equal amounts ($0.25A^\circ$) from, and to opposite sides of, the unique mean plane through the five membered ring. This conformation is confirmed by the respective observed displacements, 0.212 and $-0.209A^\circ$, from the mean plane through C(11), C(12), O(1).

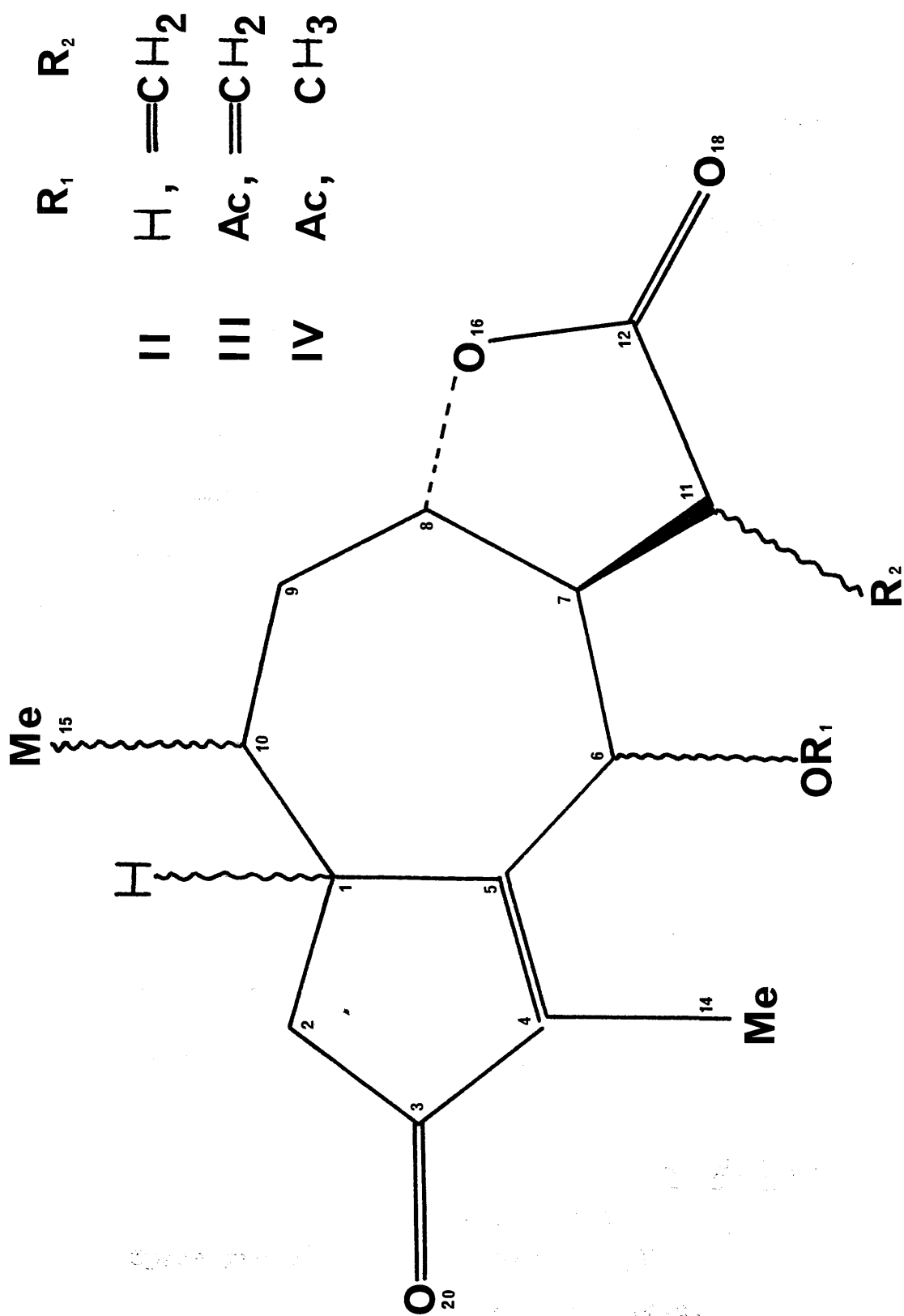
In part II of this thesis there is a further discussion on the stereochemical requirements of δ -lactones on the basis of strain energy calculations.

2.6 Mikanokryptin

Introduction

Mikanokryptin (II) has been extracted from a member of the Mikania species growing in the Canal Zone of Central America (29). The sample used in this analysis was supplied by Professor Herz (Florida State University) who has recently published (29) the results of his structure determination based mainly on spectral observations. Unambiguous assignment of the complete stereochemistry was not possible although comparison of the UV, IR and NMR spectra of related compounds allowed the main structural features to be deduced. The lactone ring

Fig. 2.6a Mikanokryptin and related compounds



fusion was elucidated by

(a) application of Samek's rule (30):

$J_{7,13}$ (translactone) $\gg J_{7,13}$ (cislactone)
to the measured chemical shifts of $J_{7,13a} =$
3.5Hz and $J_{7,13b} = 3.1\text{Hz}$ (13a and 13b represent
the two hydrogens attached to C13 (Fig. 2.6a) .

(b) the positive sign of the **C**otton effect at
257nm (3) in the **CD** spectrum of acetyl —
mikanokryptin (III).

(c) comparison of the physical properties of
acetyl dihydromikanokryptin (IV) with known
cis-fused lactones.

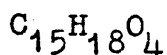
The results of these tests confirmed the trans-
fusion of the lactone ring.

The stereochemistry about C(1), C(6) and C(10)
remained uncertain and the following x-ray analysis was
carried out to determine the complete geometry of the
molecule.

2.7 Experimental

Crystal Data

Mikanokryptin

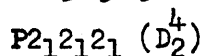


Unit cell dimensions $a = 10.366$

$$b = 9.479 \quad \alpha = \beta = \gamma = 90^\circ$$

$$c = 13.329$$

Space group



$$M = 230 \text{ a.m.u.}$$

$$V = 1301 \text{ \AA}^3$$

$$D_o = 1.151 \text{ gm cm}^{-3}$$

$$D_c = 1.176 \text{ gm cm}^{-3}$$

$$\mu = 1.03 \text{ cm}^{-1} (\text{Mo-K}\alpha_1)$$

$$F(000) = 560$$

$$Z = 4$$

The systematic absences were:

$$h00 \text{ when } h=2n+1, 0k0 \text{ when } k=2n+1, 00l \text{ when } l=2n+1.$$

These defined the space group unambiguously

The crystal used had dimensions 1 x 1 x 1.5 mm.

Data Collection

Cell dimensions and the space group were initially determined from precession photographs. The crystal was then mounted on the goniometer head of the Hilger diffractometer and aligned as described for costunolide. The revised cell dimensions are given in the crystal data. Zirconium filtered Mo-K α radiation was used to make intensity measurements by means of the θ , 2θ scan procedure.

The following scan parameters being employed:

$$\text{Increment in } \theta = 0.02^\circ$$

$$\text{No. of scan steps} = 40$$

$$\text{Time per step} = 1 \text{ sec.}$$

Immediately before and after each scan the background count was recorded for 20 seconds. Two strong reflexions were monitored as standards during data collection. These standards were checked after every 40 intensity

measurements and showed no significant variation. The intensities of the $\bar{h}\bar{k}\bar{l}$ octant were collected for $2\theta \leq 64^\circ$ and placed on a common scale by use of the standards. Integrated intensities (I) were obtained by allowing for the background as described previously. Data for which $I \leq 2\sigma I$ were not included in the subsequent calculations which yielded 2277 independent structure amplitudes.

The data in the range $2\theta \leq 20^\circ$ were also collected with reduced beam intensity so that allowance could be made for counter errors.

Structure analysis

The structure was determined by direct phasing using the multiresolution program MULTAN. Normalised structure factors were derived and triplet relationships generated for the 400 reflexions with $|E| > 1.29$. By accepting phases with a probability in excess of 0.95^s and subsequent convergence mapping the program selected three reflexions to define the origin and enantiomorph.

A further three general reflexions were chosen and the combinations of phases listed in Table XII were used to phase the 400 reflexions. An E-map was calculated from the phase set with the highest combined figure of merit. This E-map revealed all the non-hydrogen atoms.

An initial structure factor calculation gave an R of 35.5% and subsequent least squares calculations using

TABLE XII SUMMARY OF STRUCTURE DETERMINATION

A. INITIAL SET OF PHASES

REFLECTION	PHASE #	MOD. E
6 1 8 +	45 OR 315	2.51
4 4 3 +	45 OR 315	2.41
3 2 2 +	45	2.32
2 4 9	45, 135, 225 OR 315	2.18
5 0 9	90 OR 270	1.80
0 2 6	0 OR 180	1.70

B. STATISTICS

	E1	E2	E3	E4	E5	E6
THEORETICAL FOR CENTRIC	0.798	1.00	0.968	31.73	4.55	0.27
THEORETICAL FOR ACENTRIC	0.886	1.00	0.736	36.79	1.83	0.01
FOUND	0.900	1.00	0.711	34.02	1.54	0.00

* IN DEGREES
+ ORIGIN AND ENANTIOMORPH DEFINING

MOD. E MODULUS OF E FOR REFLECTION CONCERNED

E1 MEAN VALUE FOR MODULUS OF E
E2 MEAN VALUE FOR MODULUS OF E**2
E3 MEAN VALUE FOR MODULUS OF ((E**2)-1)
E4 % OF REFLECTIONS WITH MODULUS OF E > 1.0
E5 % OF REFLECTIONS WITH MODULUS OF E > 2.0
E6 % OF REFLECTIONS WITH MODULUS OF E > 3.0

the 2082 reflexions for which $|F_o| \geq 8\sigma|F_o|$ with isotropic and then anisotropic thermal parameters reduced \underline{R} to 12.1% . A difference synthesis revealed all the hydrogen atoms. Further least squares calculations with the carbon and oxygen atoms having anisotropic temperature factors and the hydrogen atoms having isotropic temperature factors yielded an \underline{R} of 4.4% . A correction was made for strong reflexions underestimated because of counter saturation, and a weighting scheme of the form:

$$W = 1/(A + BF_o + CF_o^2) \quad \text{with} \quad \begin{array}{l} A = 0.1761 \\ B = -0.0167 \\ C = 0.0018 \end{array}$$

was applied.

The values of the constants were selected so as to minimise the deviation from constant $\underline{W\Delta^2}$ over the range of $|F_o|$ and $\sin\theta$.

The refinement then converged to an \underline{R} of 4.3% . No absorption correction was applied and the ratio of observations to parameters was 8.5 to 1 .

2.8 Results

Table XIII Mikanokryptin: Fractional atomic coordinates

Positional estimated standard deviations are
shown in parentheses.

The hydrogen atoms are numbered according to the
atoms to which they are attached.

The table shows:

Atom	x	y	z
------	---	---	---

C(1)	0767(2)	5825(2)	6500(1)
C(2)	0620(2)	6477(3)	7552(1)
C(3)	-0768(2)	6264(2)	7847(1)
C(4)	-1413(2)	5479(2)	7041(1)
C(5)	-0566(2)	5204(2)	6304(1)
C(6)	-0894(2)	4321(2)	5396(1)
C(7)	0269(2)	4063(2)	4715(1)
C(8)	0746(2)	5398(2)	4191(1)
C(9)	1746(2)	6236(2)	4746(2)
C(10)	1234(2)	6921(2)	5714(1)
C(11)	0035(2)	3076(2)	3854(1)
C(12)	0775(2)	3626(2)	2988(1)
C(13)	-0657(3)	1921(3)	3812(2)
C(14)	-2794(2)	5054(3)	7116(2)
C(15)	0217(2)	8043(2)	5495(2)
O(16)	1303(2)	4884(2)	3243(1)
O(18)	0949(2)	3113(2)	2177(1)
O(20)	-1259(2)	6644(2)	8633(1)
O(21)	-1922(1)	4995(2)	4872(1)
H(1)	1429(24)	5063(27)	6509(18)
H(21)	1172(24)	6022(26)	8022(18)
H(22)	0875(29)	7530(31)	7583(22)
H(6)	-1225(21)	3440(24)	5634(17)
H(7)	0963(22)	3644(24)	5156(17)
H(8)	0037(20)	5982(23)	4010(16)
H(91)	2461(24)	5701(26)	4899(18)
H(92)	2054(25)	6984(29)	4297(19)
H(10)	1976(24)	7362(25)	6040(18)
H(131)	-0817(27)	1390(29)	3199(20)
H(132)	-1206(30)	1569(34)	4387(24)
H(141)	-3051(28)	4360(31)	6632(21)
H(142)	-3286(50)	5751(48)	7047(35)
H(143)	-3039(32)	4503(35)	7726(23)
H(151)	-0089(24)	8470(30)	6159(21)
H(152)	0543(32)	8786(33)	5115(23)
H(153)	-0531(28)	7647(31)	5110(21)
H(021)	-2270(29)	4432(30)	4540(22)

Table XIV Mikanokryptin: Thermal parameters

Carbon and oxygen values are anisotropic.

Hydrogen values are isotropic.

Estimated errors in the last two digits are shown in parentheses.

All figures are multiplied by 10^4 .

The anisotropic temperature factor expression used was of the form described by equation (12) in chapter one, the coefficients being in terms of the mean square amplitudes of vibration U_{ij} .

The table shows:

Atom	U_{11} or U	U_{22}	U_{33}	U_{12}	U_{13}	U_{23}
------	-----------------	----------	----------	----------	----------	----------

C(1)	295(8)	435(10)	281(8)	-55(8)	-18(7)	-9(8)
C(2)	431(12)	592(14)	334(9)	-119(12)	-38(9)	-94(10)
C(3)	450(11)	363(9)	307(9)	-29(10)	-17(9)	-41(8)
C(4)	352(9)	335(9)	289(8)	-47(8)	20(7)	-41(7)
C(5)	322(8)	297(8)	257(7)	-42(7)	-12(7)	4(6)
C(6)	324(8)	318(8)	251(7)	-33(7)	23(7)	-17(7)
C(7)	339(9)	324(9)	256(8)	41(7)	17(7)	9(7)
C(8)	336(9)	378(9)	289(9)	40(8)	45(7)	52(7)
C(9)	311(9)	453(11)	408(10)	-29(9)	53(9)	72(9)
C(10)	317(9)	382(10)	407(10)	-79(8)	5(8)	5(9)
C(11)	442(11)	397(10)	287(9)	92(9)	8(8)	-34(8)
C(12)	465(12)	506(13)	334(10)	167(11)	39(9)	5(10)
C(13)	734(18)	506(14)	510(14)	-79(14)	80(14)	-181(12)
C(14)	386(11)	626(15)	406(11)	-123(12)	104(10)	-105(12)
C(15)	514(14)	365(11)	610(15)	-6(11)	39(12)	6(11)
O(16)	536(9)	546(9)	314(7)	57(9)	137(7)	50(7)
O(18)	754(13)	754(13)	349(8)	145(12)	128(9)	-94(9)
O(20)	639(11)	652(11)	386(9)	-93(12)	101(8)	-218(8)
O(21)	334(7)	556(9)	386(8)	31(8)	-67(6)	-80(7)
H(1)	125(48)					
H(21)	184(61)					
H(22)	279(76)					
H(6)	84(46)					
H(7)	91(49)					
H(8)	65(43)					
H(91)	154(48)					
H(92)	218(50)					
H(10)	158(48)					
H(131)	286(72)					
H(132)	398(90)					
H(141)	290(76)					
H(142)	803(121)					
H(143)	419(73)					
H(151)	269(60)					
H(152)	360(74)					
H(153)	262(66)					
H(021)	287(73)					

TABLE XV

MIKANOKRYPIIN: INTRAMOLECULAR BONDED DISTANCES (ANGSTROMS)
STANDARD DEVIATIONS (ANGSTROMS) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	DIST.	ATOM1	ATOM2	DIST.
C1	C2	1.528(2)	C12	O16	1.353(2)
C1	C5	1.524(2)	C12	O18	1.193(2)
C1	C10	1.550(2)	C1	H1	0.998(23)
C2	C3	1.506(2)	C2	H21	0.947(22)
C3	C4	1.461(2)	C2	H22	1.037(30)
C3	O20	1.213(2)	C6	H6	0.957(21)
C4	C5	1.338(2)	C7	H7	1.009(19)
C4	C14	1.491(2)	C8	H8	0.950(19)
C5	C6	1.504(1)	C9	H91	0.921(21)
C6	C7	1.523(2)	C9	H92	0.980(22)
C6	O21	1.423(2)	C10	H10	0.978(20)
C7	C8	1.526(2)	C13	H131	0.969(24)
C7	C11	1.495(2)	C13	H132	1.006(27)
C8	C9	1.499(2)	C14	H141	0.956(27)
C8	O16	1.466(1)	C14	H142	0.837(42)
C9	C10	1.528(2)	C14	H143	0.997(24)
C10	C15	1.526(3)	C15	H151	1.019(21)
C11	C12	1.475(2)	C15	H152	0.929(28)
C11	C13	1.312(3)	C15	H153	1.001(26)

TABLE XVI MIKANOKRYPIIN: INTERMOLECULAR CONTACTS OF LESS THAN 4.0
ANGSTROMS

ATOM1	ATOM2	DIST #	ATOM1	ATOM2	DIST #
C(3).....O(21)	*	3.800	O(20).....O(21)	*	2.949
C(14).....C(11)	*	3.729	C(11).....C(14)	**	3.942
C(14).....C(12)	*	3.530	C(12).....C(14)	**	3.793
C(14).....C(13)	*	3.988	C(13).....C(14)	**	3.720
C(14).....O(16)	*	3.935	O(18).....C(14)	**	3.406
C(14).....O(18)	*	3.704	C(9).....C(15)	***	3.677
C(14).....O(21)	*	3.686	C(9).....O(21)	***	3.863
O(20).....C(6)	*	3.881	C(10).....O(21)	***	3.579
O(20).....C(11)	*	3.934	C(15).....O(21)	***	3.534
O(20).....C(13)	*	3.482			

IN ANGSTROMS

THE ASTERISKS(*) REFER TO THE FOLLOWING TRANSFORMATIONS OF THE ATOMIC
COORDINATES

* 1.5-X, 1.0-Y, 0.5+Z
 ** 0.5+X, 0.5-Y, 1.0-Z
 *** 0.5+X, 1.5-Y, 1.0-Z

TABLE XVII

MIKANOKRYPIIN: VALENCY ANGLES(DEGREES)
STANDARD DEVIATIONS(DEGREES) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	ATOM3	ANGLE	ATOM1	ATOM2	ATOM3	ANGLE
C5	C1	C2	102.8(1)	C10	C1	C2	112.0(1)
C10	C1	C5	115.3(1)	C3	C2	C1	106.1(1)
C4	C3	C2	108.3(1)	O20	C3	C2	125.8(1)
O20	C3	C4	125.9(1)	C5	C4	C3	109.4(1)
C14	C4	C3	121.9(1)	C14	C4	C5	128.7(1)
C4	C5	C1	113.2(1)	C6	C5	C1	123.8(1)
C6	C5	C4	123.0(1)	C7	C6	C5	112.6(1)
O21	C6	C5	108.0(1)	O21	C6	C7	112.1(1)
C8	C7	C6	113.1(1)	C11	C7	C6	115.1(1)
C11	C7	C8	102.9(1)	C9	C8	C7	116.1(1)
O16	C8	C7	104.0(1)	O16	C8	C9	109.0(1)
C10	C9	C8	113.3(1)	C15	C10	C1	112.2(1)
C9	C10	C1	112.9(1)	C15	C10	C9	112.2(1)
C13	C11	C7	130.1(1)	C12	C11	C7	106.7(1)
C13	C11	C12	123.2(1)	O16	C12	C11	109.2(1)
O18	C12	C11	129.3(2)	O18	C12	O16	121.5(1)
C12	O16	C8	110.3(1)	H1	C1	C2	110.5(9)
H1	C1	C5	110.2(13)	H1	C1	C10	106.0(11)
H21	C2	C1	110.9(13)	H22	C2	C1	113.6(12)
H21	C2	C3	110.3(14)	H22	C2	C3	111.1(15)
H22	C2	H21	104.9(20)	H6	C6	C7	109.8(12)
H6	C6	O21	106.4(11)	H6	C6	C5	107.7(10)
H7	C7	C8	110.6(12)	H7	C7	C11	108.3(12)
H7	C7	C6	106.8(10)	H8	C8	C9	110.5(12)
H8	C8	O16	106.4(9)	H8	C8	C7	110.3(11)

VALENCE-ANGLES (CONTINUED)

ATOM1	ATOM2	ATOM3	ANGLE	ATOM1	ATOM2	ATOM3	ANGLE
H91	C9	C8	111.8(13)	H91	C9	C10	109.3(10)
H92	C9	C10	108.4(13)	H92	C9	C8	108.1(11)
H92	C9	H91	105.6(18)	H10	C10	C15	109.2(13)
H10	C10	C1	103.6(11)	H10	C10	C9	106.2(11)
H131	C13	C11	124.2(16)	H132	C13	C11	123.6(18)
H132	C13	H131	111.6(24)	H143	C14	C4	116.0(16)
H141	C14	C4	114.1(17)	H142	C14	C4	111.4(29)
H142	C14	H141	107.6(30)	H141	C14	H141	96.2(22)
H143	C14	H142	110.3(29)	H153	C15	C10	111.7(16)
H151	C15	C10	109.2(13)	H152	C15	C10	112.3(19)
H152	C15	H151	106.2(22)	H153	C15	H151	110.3(18)
H153	C15	H152	107.0(21)	H021	O21	C6	108.4(21)

TABLE XVIII

MIKANOPYRIN: TORSION ANGLES(DEGREES)
STANDARD DEVIATIONS(DEGREES) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	ATOM3	ATOM4	ANGLE	ATOM1	ATOM2	ATOM3	ATOM4	ANGLE
C5	C1	C2	C3	-3.4(1)	C10	C1	C2	C3	121.0(1)
C2	C1	C5	C4	3.6(1)	C2	C1	C5	C6	-174.2(1)
C10	C1	C5	C4	-118.6(1)	C10	C1	C5	C6	63.6(2)
C2	C1	C10	C9	163.9(1)	C2	C1	C10	C15	-68.2(1)
C5	C1	C10	C9	-79.0(1)	C5	C1	C10	C15	48.9(2)
C1	C2	C3	C4	2.4(2)	C1	C2	C3	O20	-179.8(1)
C2	C3	C4	C5	-0.1(2)	C2	C3	C4	C14	178.0(1)
O20	C3	C4	C5	-178.0(1)	O20	C3	C4	C14	0.1(2)
C3	C4	C5	C1	-2.3(1)	C3	C4	C5	C6	175.5(1)
C14	C4	C5	C1	179.8(1)	C14	C4	C5	C6	-2.4(2)
C1	C5	C6	C7	2.5(2)	C1	C5	C6	O21	-121.9(1)
C4	C5	C6	C7	-175.1(1)	C4	C5	C6	O21	60.5(1)
C5	C6	C7	C8	-67.8(1)	C5	C6	C7	C11	174.3(1)
O21	C6	C7	C8	54.2(1)	O21	C6	C7	C11	-63.7(1)
C6	C7	C8	C9	89.9(1)	C6	C7	C8	O16	-150.4(1)
C11	C7	C8	C9	-145.3(1)	C11	C7	C8	O16	-25.6(1)
C6	C7	C11	C12	142.9(1)	C6	C7	C11	C13	-37.9(2)
C8	C7	C11	C12	19.4(1)	C8	C7	C11	C13	-161.4(2)
C7	C8	C9	C10	-67.4(1)	C7	C8	C9	C10	175.6(1)
C7	C8	O16	C12	24.0(1)	C9	C8	O16	C12	148.4(1)
C8	C9	C10	C1	62.2(1)	C8	C9	C10	C15	-65.7(2)
C7	C11	C12	O16	-5.5(2)	C7	C11	C12	O18	173.4(2)
C13	C11	C12	O16	175.2(1)	C13	C11	C12	O18	-5.8(3)
C11	C12	O16	C8	-11.9(2)	C11	C12	O16	C8	169.1(1)
C5	C1	C2	H21	116.4(15)	C5	C1	C2	H22	-125.8(16)
C10	C1	C2	H21	-119.2(15)	C10	C1	C2	H22	-1.4(16)
H1	C1	C2	C3	-121.0(12)	H1	C1	C2	H21	-1.2(19)
H1	C1	C2	H22	116.6(20)	H1	C1	C5	C4	121.5(12)
H1	C1	C5	C6	-56.4(12)	C2	C1	C10	H10	49.4(12)
C5	C1	C10	H10	166.5(12)	H1	C1	C10	C9	43.3(12)

TORSION_ANGLES (CONTINUED)

ATOM1	ATOM2	ATOM3	ATOM4	ANGLE	ATOM1	ATOM2	ATOM3	ATOM4	ANGLE
H1	C1	C10	C15	171.2(12)	H1	C1	C10	H10	-71.2(17)
H21	C2	C3	C4	-117.9(15)	H21	C2	C3	O20	60.0(15)
H22	C2	C3	C4	126.2(15)	H22	C2	C3	O20	-55.9(16)
C3	C4	C14	H141	-165.3(18)	C3	C4	C14	H142	72.5(28)
C3	C4	C14	H143	-54.8(17)	C5	C4	C14	H141	12.4(18)
C5	C4	C14	H142	-109.8(28)	C5	C4	C14	H143	122.9(17)
C1	C5	C6	H6	123.6(12)	C4	C5	C6	H6	-54.0(12)
C5	C6	C7	H7	54.1(12)	O21	C6	C7	H7	176.1(12)
H6	C6	C7	C8	172.3(12)	H6	C6	C7	C11	54.4(12)
H6	C6	C7	H7	-65.8(17)	C5	C6	O21	H021	-158.3(20)
C7	C6	O21	H021	77.1(20)	H6	C6	O21	H021	-42.9(24)
C6	C7	C8	H8	-36.7(11)	C11	C7	C8	H8	88.1(11)
H7	C7	C8	C9	-29.9(12)	H7	C7	C8	O16	89.8(12)
H7	C7	C8	H8	-156.4(17)	H7	C7	C11	C12	-97.7(12)
H7	C7	C11	C13	81.4(12)	C7	C8	C9	H91	56.6(13)
C7	C8	C9	H92	172.4(13)	O16	C8	C9	H91	-60.3(13)
O16	C8	C9	H92	55.5(13)	H8	C8	C9	C10	59.1(11)
H8	C8	C9	H91	-176.9(18)	H8	C8	C9	H92	-61.1(17)
H8	C8	O16	C12	-92.5(11)	C8	C9	C10	H10	175.1(12)
H91	C9	C10	C1	-63.2(13)	H91	C9	C10	C15	168.8(13)
H91	C9	C10	H10	49.6(18)	H92	C9	C10	C1	-177.8(13)
H92	C9	C10	C15	54.3(13)	H92	C9	C10	H10	-65.0(18)
C1	C10	C15	H151	52.1(13)	C1	C10	C15	H152	169.6(18)
C1	C10	C15	H153	-70.2(15)	C9	C10	C15	H151	-179.7(13)
C9	C10	C15	H152	-62.2(18)	C9	C10	C15	H153	58.0(15)
H10	C10	C15	H151	-62.2(18)	H10	C10	C15	H152	55.3(22)
H10	C10	C15	H153	175.5(20)	C7	C11	C13	H131	174.2(19)
C7	C11	C13	H132	4.6(20)	C12	C11	C13	H131	-6.8(19)

THE ANGLE 1-2-3-4 IS DEFINED AS POSITIVE IF, WHEN VIEWED ALONG THE 2-3 BOND, ATOM 1 HAS TO BE ROTATED CLOCKWISE TO ECLIPSE ATOM 4

TABLE XIX MIKANOKRYPTIN: MEAN PLANE CALCULATIONS

	ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
1.	C(1) C(2) C(3) C(4)	C(5) C(14) O(20)	-0.007(2) 0.012(3) -0.012(2) -0.008(2) 0.041(2) 0.035(3) -0.010(2)
2.	C(1) C(2) C(3) C(4) C(5)		-0.020(2) 0.017(3) -0.008(2) -0.006(2) 0.017(2) 0.015(3) 0.006(2)
3.	C(2) C(3) C(4)	C(14) O(20)	0.000(2) 0.000(2) 0.000(3) 0.033(2)
4.	C(3) C(4) C(5)	C(14)	0.000(2) 0.000(2) 0.000(2) -0.043(3)

TABLE XIX (CONTINUED)

	ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
5.	C(4) C(5) C(6)	C(14)	0.000(2) 0.000(2) 0.000(2) 0.048(3)
6.	C(1) C(5) C(4)	C(14)	0.000(2) 0.000(2) 0.000(2) -0.003(3)
7.	C(11) C(12) O(16)	C(7) C(8) C(13) O(18)	0.000(2) 0.000(2) 0.000(2) 0.139(2) -0.282(2) -0.092(3) 0.018(2)
8.	C(7) C(11) C(12)	C(8) C(13) O(16) O(18)	0.000(2) 0.000(2) 0.000(2) 0.495(2) -0.015(3) 0.124(2) -0.107(2)

TABLE XIX (CONTINUED)

	ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
9.	C(7) C(11) C(12) O(16)	C(8) C(13) O(18)	-0.018(2) 0.029(2) -0.031(2) 0.020(2) 0.394(2) 0.091(3) -0.128(2)
10.	C(11) C(12) O(16) O(18)		-0.002(2) 0.006(2) -0.002(2) -0.002(2) -0.153(2) 0.269(2) 0.095(3)
11.	C(13) H(131) H(132)	C(11)	0.000(3) 0.000(28) 0.000(32) 0.177(2)

Table XIX (continued)

The out of plane atoms were not included in the derivation of the planes. The equations of these planes are:

1. $0.2548x - 0.8689y + 0.4244z = 1.7314$
2. $0.2578x - 0.8627y + 0.4351z = 1.9044$
3. $0.2427x - 0.8616y + 0.4458z = 1.8698$
4. $-0.2430x + 0.8616y - 0.4456z = -1.8707$
5. $-0.2982x + 0.8224y - 0.4846z = -2.9317$
6. $0.2708x - 0.8426y + 0.4655z = 2.4037$
7. $0.8182x - 0.4698y + 0.3314z = 8.8434$
8. $-0.7901x + 0.5461y - 0.2784z = -8.0568$
9. $-0.8239x + 0.4947y - 0.2716z = -8.5435$
10. $-0.8157x + 0.4657y - 0.3389z = -8.8849$
11. $-0.7198x + 0.6464y - 0.2592z = -7.0786$

2.9 Discussion of results

Tables XIII - XIX⁺⁺ summarise the results of the x-ray analysis. A view of the molecule is given in Fig. 2.8 and the crystal packing viewed along the b axis is illustrated

⁺⁺ The torsion angles in Table XVIII give the enantiomer and the signs should therefore be reversed to obtain the correct absolute stereochemistry.

Fig. 2.8 Mikanokryptin; A view of the molecule



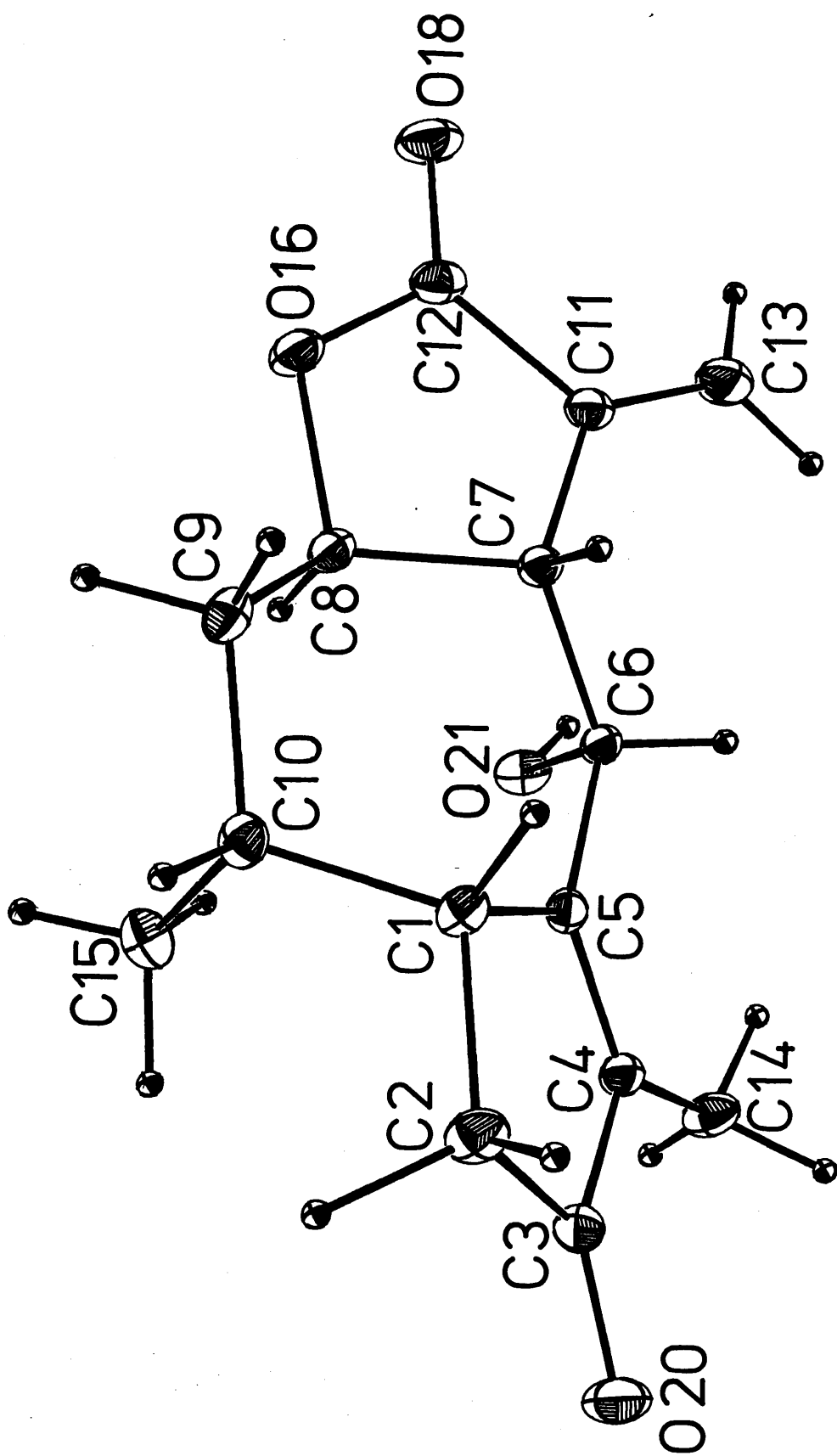
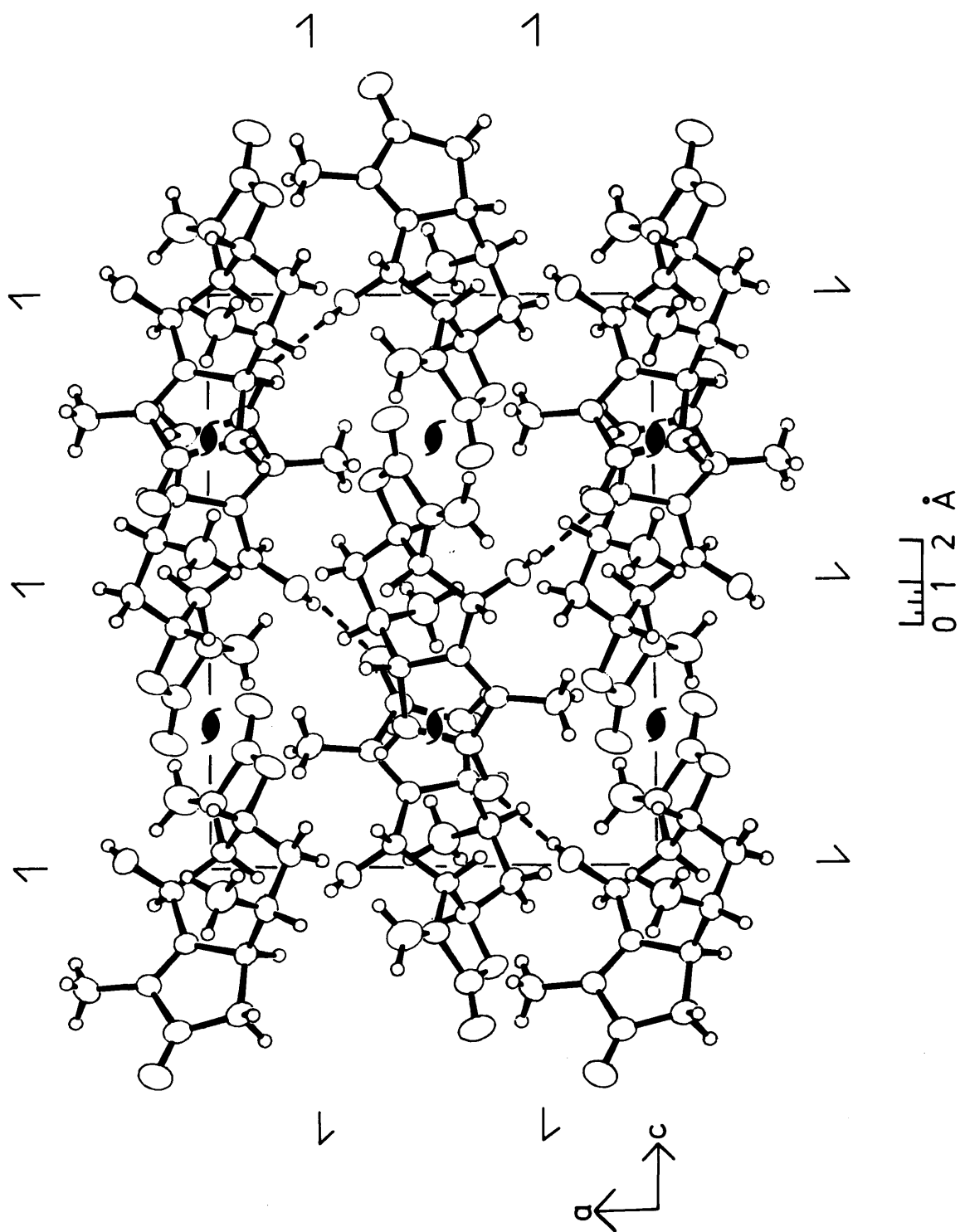




Fig. 2.9 Mikanokryptin crystal packing: A view down the 'b' axis





in Fig. 2.9 . The final and observed structure factors are listed in Table XXII.

The stereochemistry of mikanokryptin was established as shown in Fig. 2.10 . The absolute configuration was not determined, but if it is assumed that the C(7) side chain is β , as in almost all sesquiterpene lactones*, then the relative orientations of the asymmetric centres are in agreement with the predictions of Herz et al (29). That is, for example, H(1) α , C(6) hydroxyl β , and C(10) methyl β ,

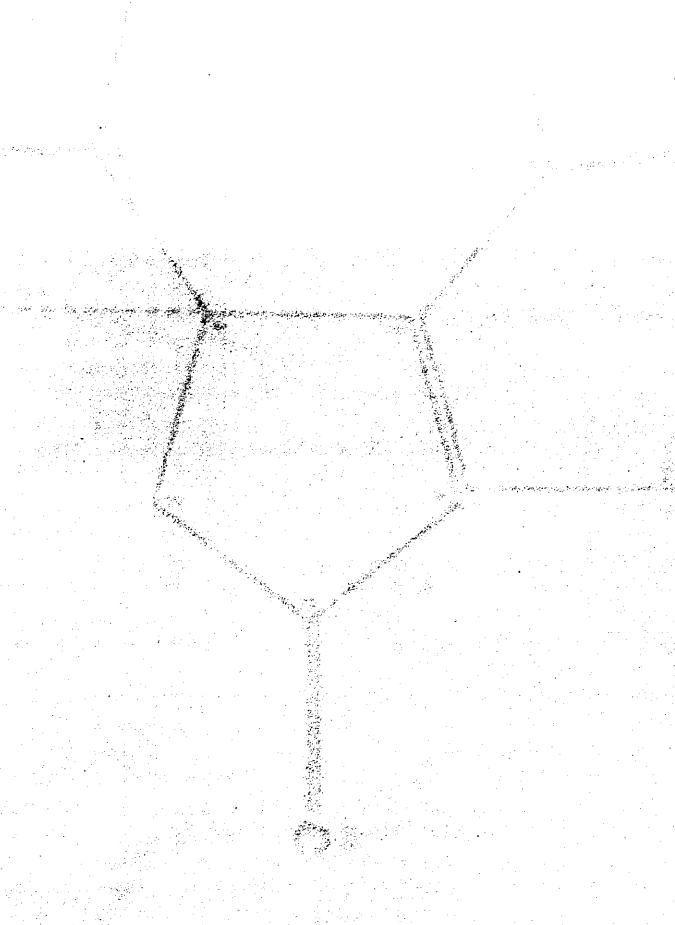
Intermolecular hydrogen bonding occurs between the carbonyl O(20) in one molecule and the C(6) hydroxyl function in another (Table XVI). The oxygen atoms are separated by 2.95⁹Å and the H(O21)....O(20) distance is 2.20⁹Å. These three atoms are situated so as to form an O(20)....H(O21) - O(21) angle of 162° .

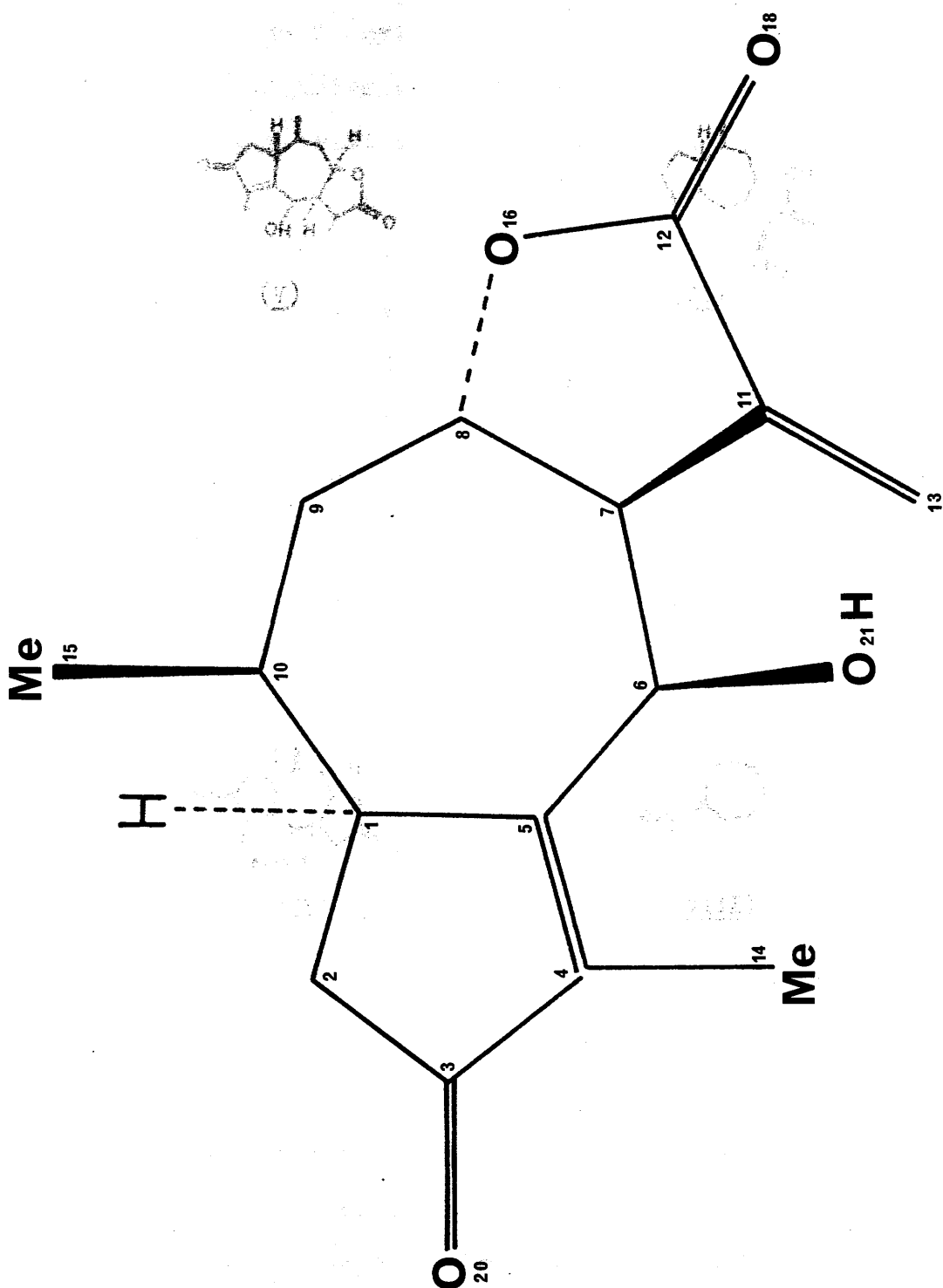
Mikanokryptin has the unusual structural feature of an sp² carbon atom at one of the points of fusion of the five and seven membered carbocycles and only a small number of similar guianolides has been studied. Both geigerin (V) and some of its analogues (31,32) and aciphyllic acid (VI) (33) have been examined by spectral techniques, while the only corresponding crystal structure analysis in the literature is the report (34) on bromogeigerin (VII).

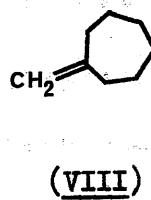
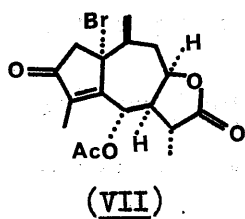
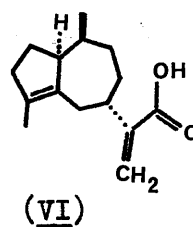
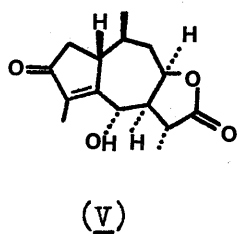
However, calculations (35,36,37,38) have been carried out for isolated cycloheptanes and cycloheptenes and x-ray

* An exception is Centurepensin, a naturally occurring guianolide containing chlorine. See: A. T. Hewson, R. C. Pettersen, and O. Kennard, Cryst. Struct. Comm; 1, 383, (1972) .

Fig. 2.10 Structure of Mikanokryptin



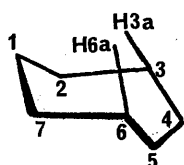




results are available for some guianolides (e.g. 39,40,41) containing these systems.

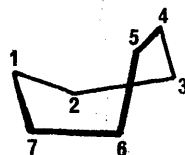
It is well known that both cycloheptane and cycloheptene can exist in families of chairs and boats (Fig. 2.11). In the cycloalkane there is a C_2 and C_s form within each family, with the former being the most stable. In the C_s conformers, chairs are preferred to boats by 1.0 to 2.5 K.cal(35). The twist chairs are more stable than the chairs mainly because of a strong H(3a)....H(6a) transannular interaction in the latter (Fig. 2.11), which is relieved by pseudorotation to a twist form. The cycloalkene presents a slightly different situation in that the two sp^2 carbon atoms, with normal valence angles of 120° , facilitate an increase in the H(3a)H(6a) separation and most workers agree (36,37,38) that the C_2 twist conformations are just slightly less stable than those possessing a plane of symmetry. This is supported by the fact that spectroscopic studies have found a preference for cycloheptene to exist in the chair form (42,43).

The cycloheptane ring in mikanokryptin represents a situation somewhere between cycloheptane and cycloheptene and as a suitable force field was available (44, Appendix A), it was decided to calculate minimum energies and geometries for the twist chair and chair for 1-methyldene cycloheptane (VIII). The boat family was not included in the calculation since this is generally accepted to be less stable. The force field was also used for calculations of the C_2 and C_s chairs of cycloheptane and cycloheptene so that these results could be correlated with the data already published (Table XX).

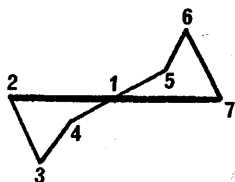


Chair

\underline{C}_s

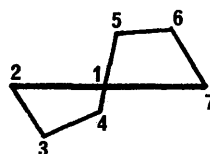


Boat



Twist-chair

\underline{C}_2



Twist-boat

Similar conformations exist for cycloheptene

Fig. 2.11 Major conformers of cycloheptane

TABLE XX CONFORMATIONS AND ENERGIES OF VARIOUS SEVEN-MEMBERED RINGS

OMEGA	CYCLOHEPTANE				CYCLOHEPTENE				VIII			
	CHAIR HE	CHAIR TH	T. CHAIR HE	T. CHAIR TH	CHAIR ER	CHAIR TH	T. CHAIR ER	T. CHAIR TH	CH TH	TC TH	MK CR	
1	-70	-65	-76	-74	-68	-70	-28	-32	-65	-70	-67	
2	92	85	97	86	75	75	78	79	84	86	90	
3	-71	-67	-41	-38	-59	-58	-43	-43	-67	-44	-68	
4	0	0	-41	-38	0	0	-43	-43	1	-33	3	
5	71	67	97	86	59	58	78	79	65	84	63	
6	-92	-85	-76	-74	-75	-75	-28	-32	-84	-77	-79	
7	70	65	53	58	68	70	-8	-2	66	59	62	
SIG2	114	105	0	0	92	90	0	0	108	19	104	
SIGS	0	0	141	114	0	0	154	160	4	93	24	
QUOT	1.00	1.00	---	---	1.00	1.00	---	---	0.96	0.17	0.81	
ENERGY												
HE	2.16		0.00									
ER					0.00		0.42					
FA					0.00		1.71					
AL					0.00		0.57					
TH	0.74		0.00		0.00		1.53		0.00	0.94		

ENERGIES (K.CAL/MOLE) ARE QUOTED RELATIVE TO THE MOST STABLE ISOMER IN EACH SET OF CALCULATIONS

SYMBOLS EXPLAINED ON FOLLOWING PAGE

EXPLANATION OF SYMBOLS IN TABLE AX

HENDRICKSON(35)
 ERMER(38)
 FAVINI(36)
 ALLINGER(37)
 CALCULATIONS REPORTED IN THIS THESIS

TORSION ANGLES IN SEVEN MEMBERED RING AS SHOWN IN FIG.2.12
 SEE FIG.2.12 FOR EXPLANATION
 SEE FIG.2.12 FOR EXPLANATION
 SEE FIG.2.12 FOR EXPLANATION

CHAIR CONFORMATION
 TWIST CHAIR CONFORMATION
 TWIST CHAIR CONFORMATION
 MIKANOKRYPTIN CRYSTAL STRUCTURE

OMEGA
 SIG2
 SIGS
 QUOT

CH
 T-CHAIR
 TC
 MK/CR

These results confirm the relative values obtained by Hendrickson (35) for the chair and twist chair of cycloheptane, the latter conformation being more stable by 0.74 K.cal per mole (Hendrickson 2.16 K.cal per mole).

In cycloheptene the calculations of Favini (36), Allinger (37), and Ermer (38) predicted a preference for the C_s over the C₂ form and a similar relationship has been produced in this work. Although the enthalpy difference of 1.5 K.cal per mole is somewhat larger than those of Allinger (0.57) and Ermer (0.42) the geometry is very similar to that computed by Ermer, and the agreement with the experimental data, that, for most derivatives the chair is preferred by about 1-2 K.cal per mole (45), is good.

Thus the calculated values for the geometries and energies of cycloheptane and cycloheptene are consistent with those of other workers. Similar calculations for VIII indicated that the most stable conformation corresponds more closely to C_s than C₂ symmetry.

In mikanokryptin the carbocycle corresponding to VIII adopts a chair conformation in contrast to the more frequently observed twist chair in the cycloheptane rings of guianolides. The chair conformation is most common when the guianolide structure contains a cycloheptene system as shown in Table XXI.

In this table the torsion angles ω_1 to ω_7 are arranged so that the C_s plane in the chair conformations and the C₂

TABLE XXI CONFORMATION OF THE SEVEN MEMBERED RING IN VARIOUS GUANOLIDES

CYCLOHEPTANES					==#	CYCLOHEPTENES							
OMEGA	EB	BS	BH	BM	UH	SO	IF	BG	MK	BL	UG	BL	CL
1	-49	-72	61	51	-76	-69	-59	-69	67	-93	-66	-57	33
2	85	81	-98	-86	87	92	80	90	-90	69	76	76	-76
3	-42	-42	53	60	-51	-45	-68	-43	68	-30	-61	-73	28
4	-41	-31	32	19	-23	-35	3	-37	-3	-1	4	6	49
5	102	79	-81	-82	65	88	68	84	-63	29	60	64	-78
6	-63	-78	59	68	-64	-74	-87	-59	79	-52	-85	-82	22
7	25	63	36	-36	65	51	62	44	-62	74	69	64	11
FUSN	CT*	CT*	TC	CC	TC	CT*	TC	-C	-T	-T	TT	TT*	-C
SIG2	32	19	40	62	62	19	101	22	104	110	92	116	34
SIG5	147	80	154	74	71	114	13	134	24	38	17	28	197
QUOT	0.18	0.19	0.21	0.46	0.47	0.14	0.89	0.14	0.81	0.74	0.84	0.80	0.17

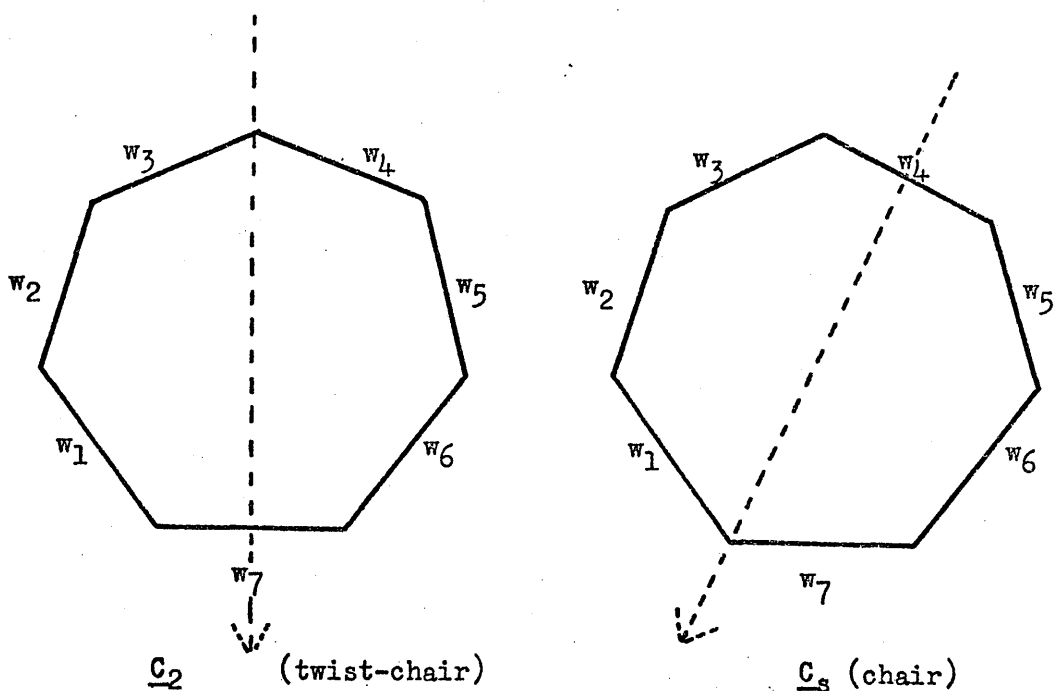
SYMBOLS EXPLAINED ON FOLLOWING PAGE

EXPLANATION OF SYMBOLS IN TABLE XXI

EB	EUPAROTIN BROMOACETATE (40)
BS	BROMODIHYDROISOPHTOSANTONIC LACTONE ACETATE (46)
BH	BROMOHELANIN (47)
BM	BROMOMEXICANIN-E (48)
OH	AN OXIDE OF HELANIN (49)
SO	SULISTAN (50)
IF	4-ACETYL-2-P-IOUBENZOYLFLORILENALIN (51)
BG	BROMOGEIGERIN (34)
MK	MIKANOKRYPTIN (TORSION ANGLES HAVE OPPOSITE SIGNS TO THOSE IN TABLE XVIII, TO GIVE CORRECT ABSOLUTE CONFIGURATION)
BL	BROMOGAILLANDIN (39)
DG	DEACETYLDIHYDROGAILLANDIN P-BROMOBENZOATE (39)
BL	BERLANDIN (41)
CL	CAROLENALONE (52)
==#	SEVEN MEMBERED RING INCORPORATES AN EXOCYCLIC DOUBLE BOND WHICH IS NOT LOCATED AT ONE OF THE POINTS OF FUSION TO THE CYCLOPENTANE/ENE
==#	SEVEN MEMBERED RING INCORPORATES AN EXOCYCLIC DOUBLE BOND WHICH IS LOCATED AT ONE OF THE POINTS OF FUSION TO THE CYCLOPENTANE/ENE
OMEGA	TORSION ANGLES IN SEVEN MEMBERED RING AS SHOWN IN FIG. 2.12
FUSN	STEREOCHEMISTRY OF RING FUSIONS IN THE ORDER CYCLOPENTANE/ENE, LACTONE. THE SYMBOLS C, T REPRESENT CIS, TRANS RESPECTIVELY. THE SYMBOL -- REPRESENTS A FUSION IN WHICH THERE IS AN (SP**2) CARBON ATOM AT ONE OF THE RING JUNCTIONS. THESE LACTONES ARE D, 7 FUSED. ALL OTHERS ARE 7.8
*	SEE FIG. 2.12 FOR EXPLANATION
SIG2	SEE FIG. 2.12 FOR EXPLANATION
SIGS	SEE FIG. 2.12 FOR EXPLANATION
QUOT	SEE FIG. 2.12 FOR EXPLANATION

axis in the twist forms are located as described by McPhail and Sim (40) and illustrated in Fig. 2.12 which also gives their algorithm for calculating the deviations from absolute C_s or C_2 symmetry.

Table XXI shows quite clearly that there is a wide range of possible conformations in the C-7 ring of guianolides. Only the chair and twist chair variants have been considered in this compilation, but boat-like stereochemistry is also known (53). The conformation adopted is probably influenced by the requirements of the modes of fusion of the two five membered rings. In compounds I to V in Table XXI for instance three different permutations are apparent. Euparotin bromoacetate (EB) and bromodihydroisophotosantonin lactone acetate (BS) have almost identical values of the function $\Sigma_2/(\Sigma_2+\Sigma_s)$ and thus both seven membered rings are closer to twist chairs than chairs although the chair form of cycloheptane would apparently require less distortion to produce the stereochemistry observed in the vicinity of the ring fusions than the twist chair (Fig. 2.13). However, the calculations have shown that for an isolated cycloheptane the C_2 form is more stable and because of the predominance of this conformation in the guianolides incorporating a cycloheptane ring it would appear that the minimum energy conformation of the isolated ring is adopted if at all possible. The torsion angles in this ring for euparotin bromoacetate correspond most closely to the twist chair and Figure (2.13) illustrates that the calculated minimum energy twist chair would permit cis fusion of the cyclopentene ring (with an observed C4-C5-C1-C2 angle of $+18^\circ$) more readily than the corresponding trans fusion and the



w_1 to w_7 are the respective endocyclic torsion angles

$$\Sigma_2 = (|w_1 - w_6| + |w_2 - w_5| + |w_3 - w_4|)$$

$$\Sigma_s = (|w_1 + w_7| + |w_2 + w_6| + |w_3 + w_5| + |w_4|)$$

$$\text{QUOT} = \Sigma_2 / (\Sigma_2 + \Sigma_s)$$

QUOT

approaches 1.0 for conformations close to chair

approaches 0.0 for conformations close to twist-chair

Fig. 2.12 Relationships between cycloheptane torsion angles and conformation(40)

Fig. 2.13 Euparotin Bromoacetate: Stereochemistry in the vicinity of the ring fusions.

(a) - (d) Newman projections down the C(5)-C(1) and C(6)-C(7) bonds of the calculated chair and twist-chair of cycloheptane (FFI).

(a) Twist-chair down C(5)-C(1)

(b) Chair down C(5)-C(1)

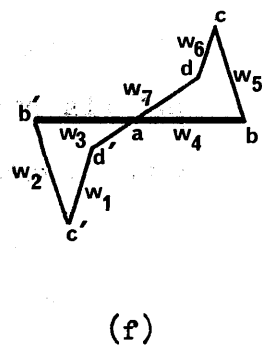
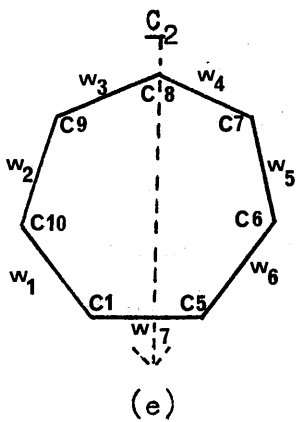
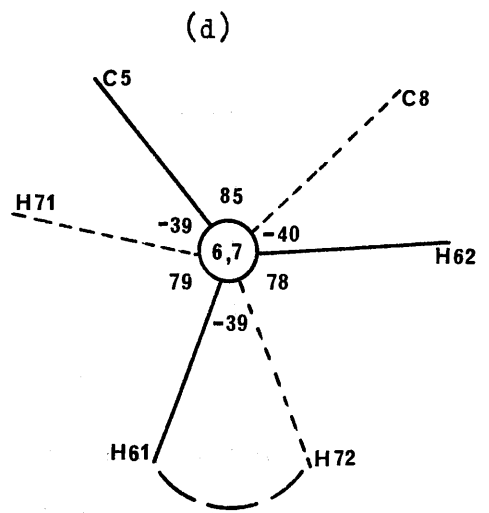
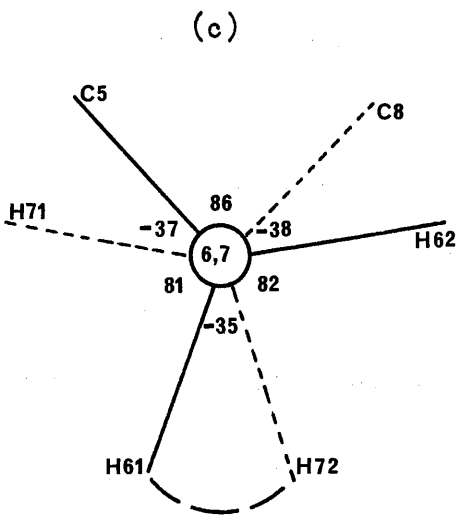
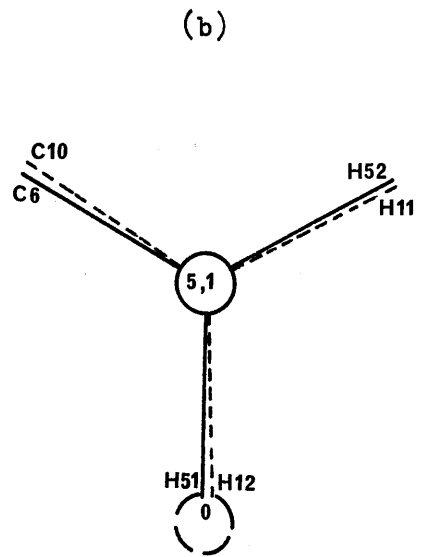
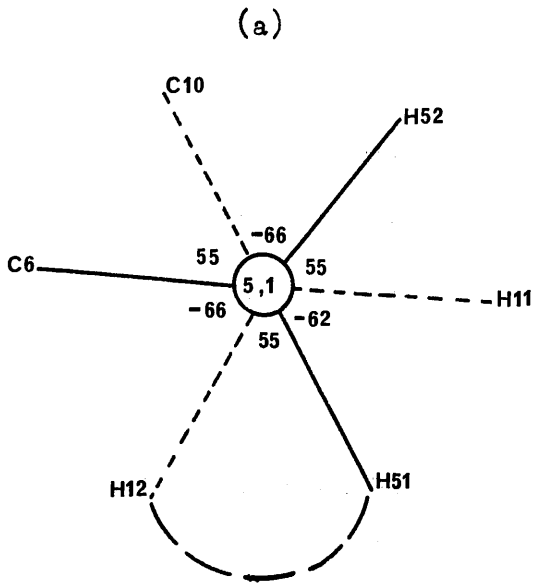
(c) Twist-chair down C(6)-C(7)

(d) Chair down C(6)-C(7)

Torsion angles are quoted in degrees and the (longer) dashed lines indicate the points of fusion which would produce the correct stereochemistry.

(e) Location of the C_2 axis in the crystal structure of this compound and relationship to the torsion angles (w_1 to w_7) used in Tables XX and XXI.

(f) Correlation between the locations preferred for fusion to an isolated cycloheptane as reported by Hendrickson(35) and the torsion angles w_1 to w_7 .



formation of a C(7) β -fused- γ -lactone is also relatively strain free. This is in agreement with the prediction of Hendrickson(35) and the cyclopentene and lactone rings are fused at the d-d' and b-c bonds respectively (Fig. 2.13).

The related compounds bromohelamin and the oxide of helamin (Table XXI) provide an example of the change in conformation of the cycloheptane in order to accommodate the requirements of the gross structure.

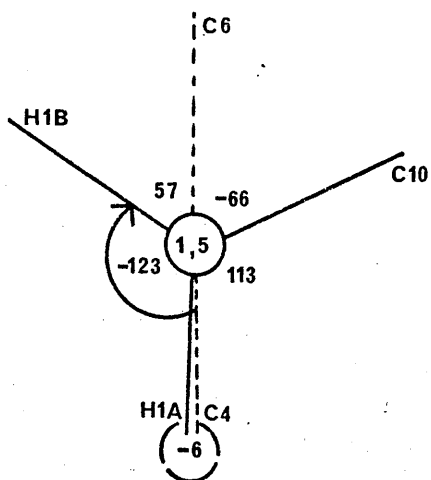
Although both these molecules have identical modes of fusion of the smaller rings, the latter compound has a cycloheptane ring which has adopted a conformation somewhere between the twist chair and chair, presumably in order to be able to form the oxygen bridge between the five and seven membered carbocycles.

The conformation of the cycloheptane ring in mikanokryptin is much closer to a chair than a twist chair and Fig. 2.14 illustrates that relatively little additional strain is imposed on an isolated chair when it is fused to the smaller rings with the observed stereochemistry. Cycloheptene rings in guianolides generally adopt a twist chair conformation but again the minimum energy stereochemistry of the isolated ring may not always represent the most stable conformation in a crystal structure.

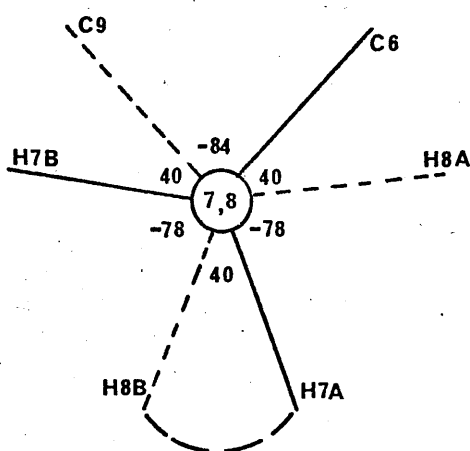
Further calculations are required to allow for the different locations of the C_2 axis or C_s plane in the non-symmetrical seven-membered rings before all the observed structures can be interpreted in terms of the stereochemistry of ring fusion.

Fig. 2.14 Mikanokryptin: Stereochemistry in the vicinity of ring fusions.

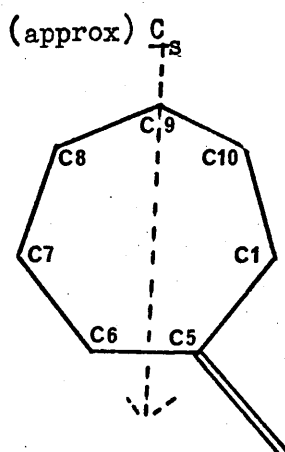
- (a) and (b) Newman projections down the C(1)-C(5) and C(7)-C(8) bonds respectively of calculated (FFI) 1-Methyldene-cycloheptane (chair conformation).
- (c) Location of the approximate C_s plane in Mikanokryptin.



(a)



(b)



(c)

The mean of the valency angles in the cycloheptane ring is 115.4° and this is very similar to that found in other guianolides (40). This infers that the average Csp^3 valence angle is less distorted in the mikanokryptin structure since this mean includes the lone Csp^2 angle thus supporting previous statements regarding the stabilising effect of this atom.

The cyclopentenone ring is fused to the seven-membered carbocycle at C(1) and C(5) with the C(4)—C(5)—C(1)—C(2) torsion angle adopting a value of 4° . Inspection of a Dreiding model of the larger ring shows that its Cs chair form would allow complete eclipsing of the C(4) - C(5) and C(1) - C(2) bonds and that increasing pseudo-rotation towards the twist chair produces greater staggering. The torsion angles in the smaller carbocycle clearly indicate that it is almost planar and this is supported both by the maximum displacement (0.02\AA for C(1)) from the mean plane through the atoms of this ring and by calculations using the algorithm of Cremer and Pople (28) which indicate a flattened envelope ($q = 0.03\text{ \AA}, \Phi = 2^\circ$). A similar though less flattened envelope has also been observed in berlandin. The mean of the endocyclic valence angles is 108° in the mikanokryptin cyclopentenone ring and 107° in the corresponding portion of the berlandin structure. These values compare very favourably with the expected angle of 108° for a planar cyclopentane.

The α -methylene - γ - lactone is transfused at C(7) - C(8) and adopts a conformation which is between the half chair with the approximate C₂ axis through the centre

of the C(7) - C(8) bond and the envelope with C(8) out of the plane of the other four atoms. In the notation of Cremer and Pople $q = 0.25\text{\AA}^\circ$ and $\phi = 23^\circ$ and this is confirmed by the fact that C(7) and C(8) are displaced, by 0.14 and 0.28\AA° respectively, to opposite sides of the plane through C(11), C(12) and O(16).

The positive Cotton effect at 257nm for acetylmikanokryptin (29) has been correlated with a C(7) β - C(8) trans-fused lactone by Stöcklin et al (3). As stated previously, the absolute configuration was not determined in this analysis and the torsion angles in Table XVIII should have their signs reversed to represent the stereochemistry predicted by the Cotton effect. Thus the C(13) - C(11) - C(12) - O(18) torsion angle becomes $+6^\circ$ which supports the predictions of Beecham (27) regarding the relationship between the sign of the Cotton effect and the chirality of this chromophore in lactones trans-fused to six and seven-membered rings.

Table XIIa Costunolide: Final structure factors

The table shows:

L^*	$ F_o $	$ F_c $	Phase($^\circ$)
-------	---------	---------	-------------------

* Reflections sorted into groups with common
H,K indices (shown in each group heading).

0,0,L	5	55	57	270	5	96	92	0	7	39	40	90	0,13,L	1	16	17	90
2 175 179 180	6	41	40	90	6	135	135	0	8	32	33	90		3	9	10	90
4 30 29 0	7	30	30	270	7	30	27	180					0,18,L				
8 12 10 180	8	15	14	90	8	15	14	180	0,10,L	1	95	93	90	0	15	16	180
	9	17	16	0				0		2	41	40	90	1	11	13	180
0,1,L										3	47	47	90	2	12	11	0
0,4,L	0	413	42	0	0	223	230	0	0	5	8	3	90				
1 179 201 90	1	380	380	180	1	65	60	0	0,14,L								
2 394 375 90	2	242	225	0	2	29	27	180		0	34	35	180	1	409	436	270
3 93 97 90	3	79	77	180	3	15	7	0		1	21	19	180	2	113	109	180
4 448 470 90	4	20	9	180	4	46	44	0		2	17	17	0	3	127	117	270
5 85 84 270	5	49	44	180	5	68	71	90		3	64	63	180	4	133	131	0
6 20 20 90	6	32	30	0	6	73	74	90	0,11,L	4	43	42	180	5	128	130	270
7 62 63 90	7	92	89	0	7	25	27	270		5	36	39	180	6	31	33	180
8 25 25 270	8	48	49	180						6	24	24	180	7	56	58	90
0,2,L																	
0,5,L	0	184	183	270	0	139	135	180	1	38	37	90	0,15,L	1	51	52	90
1 815 905 180	1	275	282	270	1	380	395	0	2	75	74	90		2	17	19	90
2 665 669 0	2	148	141	90	2	209	213	0	3	92	95	90		3	8	0	270
3 28 4 180	3	7	1	270	3	77	79	0	4	102	103	90		4	0	90	
4 149 154 0	4	88	83	270	4	83	82	0	5	19	17	90		5	8	90	
5 130 127 180	5	27	25	90	5	32	34	0	6	15	15	90		6	0	270	
6 98 98 180	6	11	12	90	6	26	25	180	0,12,L	7	9	8	90	7	281	283	157
7 39 42 0	7	15	15	90	7	10	12	180						8	49	49	99
8 73 70 0	8				8	11	13	0						5	65	58	68
0,3,L														6	26	25	148
0,6,L	0	232	239	180	0	129	129	0	0,16,L					7	20	21	51
1 705 746 90	1	281	275	180	1	47	53	180		0	30	27	0	8	28	26	265
2 769 772 90	2	82	83	0	2	21	21	180		1	22	25	180				
3 8 7 270	3	46	35	0	3	34	32	180		2	39	43	180	1	100	88	90
4 34 37 270	4	14	3	180	4	11	11	180		3	20	19	180	2	640	676	309
	5				5	31	33	180	0,17,L	4							

3,3,L	3,6,L	3,9,L	3,12,L	3,16,L	4,2,L
3 137 142 43	0 63 61 270	0 208 211 270	0 63 61 90	0 13 13 270	2 291 307 44
4 187 192 203	1 52 51 343	1 67 69 240	1 47 47 324	1 41 43 250	3 224 237 287
5 98 101 100	2 130 133 206	2 21 19 321	2 94 91 306	1 30 31 165	4 136 134 262
6 62 60 355	3 248 258 314	3 27 25 166	3 71 67 21	2 22 22 316	5 74 78 106
7 33 34 0	4 60 59 355	4 119 117 317	4 27 29 253	3 22 22 316	6 83 83 205
8 19 16 337	5 7 9 236	5 42 41 131	5 20 21 286	3 17,L	7 33 32 249
3,4,L	6 41 46 75	6 28 29 160	6 20 22 346	0 25 27 90	8 19 19 266
	7 30 32 99	7 10 9 293	7 19 21 312	0 25 26 235	
		8 22 23 224		1 25 27 90	
	3,7,L	3,10,L	3,13,L	3,18,L	
0 178 170 270	0 129 131 270	0 89 89 270	0 90 88 90	0 10 11 270	0 137 137 180
1 336 336 305	1 61 61 170	1 122 124 28	1 39 37 171	2 6 7 18	1 58 63 280
2 37 39 92	2 149 149 311	2 13 15 187	2 69 71 245	3 8 8 145	2 219 228 151
3 50 47 281	3 90 87 56	3 46 44 325	3 32 31 24	0 10 11 270	3 133 143 137
4 124 123 274	4 54 50 317	4 108 107 5	4 27 28 68	1 7 6 179	4 109 111 226
5 103 101 18	5 54 55 313	5 15 15 51	5 32 31 24	0 10 11 270	5 77 77 94
6 27 27 348	6 36 38 63	6 28 29 261	6 27 28 68	1 7 6 179	6 23 19 143
7 24 23 359	7 39 41 110	7 10 9 351	7 27 28 68	0 10 11 270	7 42 39 52
8 8 7 185	8 26 25 223	8 10 9 351	8 27 28 68	4,0,L	8 7 5 330
9 19 20 148					9 5 2 167
3,5,L	3,8,L	3,11,L	3,14,L	4,3,L	
0 141 147 270	0 195 201 270	0 140 137 270	1 51 50 355	0 338 332 180	0 112 109 180
1 112 113 250	1 57 55 335	1 130 130 78	2 16 15 147	1 214 210 90	1 113 117 121
2 142 152 23	2 198 199 344	2 42 39 292	3 20 21 236	2 110 107 180	2 224 232 222
3 63 66 181	3 31 29 20	3 56 55 212	4 18 17 269	3 97 90 90	3 73 70 225
4 60 62 26	4 105 102 104	4 22 22 214	5 23 22 314	4 107 106 180	4 71 73 110
5 163 164 6	5 68 68 17	5 35 34 145	6 26 28 300	5 9 13 90	5 67 71 278
6 83 86 254	6 50 51 50	6 11 11 5	7 8 6 90	6 93 95 180	6 36 37 295
7 45 43 42	7 21 22 282	7 14 15 183	8 19 19 214	7 55 54 90	7 46 48 225
8 25 24 71	8 11 9 55	8 14 15 183	9 19 19 214	8 13 13 180	8 30 30 218
9 9 11 92			10 12 13 342	4,1,L	
				0 330 316 180	
				1 77 72 53	

5,5,L	0	84	87	270	4	21	20	303	1	8	8	241	4	32	33	289	3	49	52	208
115 117 306	1	69	70	352	5	30	28	63	2	19	18	279	5	24	27	58	4	28	28	277
126 126 280	2	41	44	61	6	25	27	226	3	15	15	254	6	14	15	219	5	45	44	48
71 69 19	3	92	91	292	7	7	6	83	4	16	16	141	7	57	57	176	6	31	30	179
97 98 235	4	58	59	158									8	25	27	252	7	18	17	205
85 88 358	5	56	55	57													8	18	18	267
50 52 339	6	43	43	21																
11 10 317	7	16	17	109																
5,6,L																				
107 106 270	1	22	24	217	0	35	35	90	0	5	7	90	0	116	116	180	0	159	158	0
95 92 301	2	85	83	44	1	35	35	180	1	9	10	90	1	50	52	33	1	85	87	220
40 38 338	3	56	54	28	5	34	33	182	2	27	29	74	2	165	166	127	2	34	32	150
52 54 336	4	45	44	226	6	28	27	127	3	20	21	22	3	156	157	123	3	67	69	101
39 38 315	5	23	22	155				8					4	85	86	138	4	108	111	265
32 30 78	6	16	16	25					0	5	5	270	5	35	33	145	5	53	52	245
50 52 85	7	12	12	240					1	11	10	296	6	29	31	245	6	10	10	161
32 33 37													8	21	22	252	7	14	17	352
13 12 52																	8	10	10	46
5,7,L																				
5,10,L																				
33 33 74	1	33	33	74	0	41	43	90	0	17	18	180	0	151	150	0	0	105	107	180
95 94 208	2	95	94	208	1	17	17	290	1	65	64	270	1	120	120	44	1	39	39	73
22 19 212	3	22	19	212	2	25	24	265	2	51	53	180	2	99	100	286	2	71	73	155
58 58 78	4	58	58	78	3	35	36	167	3	116	115	90	3	37	38	85	3	41	37	320
21 20 79	5	30	29	9	4	14	14	167	4	113	116	180	4	53	51	164	4	54	55	112
7 8 261	6	7	8	261	5	9	10	268	5	112	114	90	5	34	33	249	5	55	56	254
5,11,L																				
12 11 90	0	12	11	90	0	46	45	240	0	11	10	180	0	109	115	210	0	14	15	162
8 8 83	1	8	5	355	1	8	5	83	1	44	42	90	1	17	17	263	1	21	20	317
5,15,L																				
32 32 330	4	32	32	330	5	9	11	80	5	7	4	180	5	5	6	9	5	13	15	311
6,1,L																				
12 14 0	0	12	14	0	0	12	14	0	0	12	14	0	0	143	140	180	0	90	88	0
187 186 57	1	187	186	57	1	187	186	57	1	187	186	57	1	120	124	187	1			
102 101 164	2	102	101	164	2	102	101	164	2	102	101	164	2	208	217	136	2			
112 117 206	3	112	117	206	3	112	117	206	3	112	117	206	3				3			
6,3,L																				
151 150 0	0	151	150	0	0	151	150	0	0	151	150	0	0	120	120	286	0	105	107	180
120 120 44	1	120	120	44	1	120	120	44	1	120	120	44	1	99	100	286	1	39	39	73
99 100 286	2	99	100	286	2	99	100	286	2	99	100	286	2	37	38	85	2	71	73	155
37 38 85	3	37	38	85	3	37	38	85	3	37	38	85	3	53	51	164	3	41	37	320
53 51 164	4	53	51	164	4	53	51	164	4	53	51	164	4	34	33	249	4	54	55	112
34 33 249	5	34	33	249	5	34	33	249	5	34	33	249	5	109	115	210	5	55	56	254
109 115 210	6	109	115	210	6	109	115	210	6	109	115	210	6	17	17	263	6	14	15	162
17 17 263	7	17	17	263	7	17	17	263	7	17	17	263	7	5	6	9	7	21	20	317
5 6 9	8	5	6	9	8	5	6	9	8	5	6	9	8				8	13	15	311
6,4,L																				
143 140 180	0	143	140	180	0	143	140	180	0	143	140	180	0				0	90	88	0
120 124 187	1	120	124	187	1	120	124	187	1	120	124	187	1				1			
208 217 136	2	208	217	136	2	208	217	136	2	208	217	136	2				2			
6,7,L																				
105 107 180	0	105	107	180	0	105	107	180	0	105	107	180	0				0	90	88	0
39 39 73	1	39	39	73	1	39	39	73	1	39	39	73	1				1			
71 73 155	2	71	73	155	2	71	73	155	2	71	73	155	2				2			
41 37 320	3	41	37	320	3	41	37	320	3	41	37	320	3				3			
54 55 112	4	54	55	112	4	54	55	112	4	54	55	112	4				4			
55 56 254	5	55	56	254	5	55	56	254	5	55	56	254	5				5			
14 15 162	6	14	15	162	6	14	15	162	6	14	15	162	6				6			
21 20 317	7	21	20	317	7	21	20	317	7	21	20	317	7				7			
13 15 311	8	13	15	311	8	13	15	311	8	13	15	311	8				8			

[illegible]

6	9,1,L	6	21	22	244	1	82	81	215	2	12	13	84	10,3,L	5	19	19	5
7	10 9 195		9,5,L			2	25	26	158	3	12	13	120					
	16 17 230	0	115 118 90			4	11 12 99	16 173			9,13,L			0 112 109 0		10,7,L		
	9,2,L	1	36 37 211			5	17 16 173			0	5	5 90		1 31 31 335	0	24	25 180	
0	92 96 90	2	108 104 118							1	16 16 329			2 15 12 145	1	8	8 221	
1	82 82 22	3	11 10 98			0	34 34 90			2	17 16 103			3 38 39 314	0	21	21 60	
2	49 50 156	4	29 31 221			1	15 17 346			0	10 11 351			4 18 18 342	1	12	13 337	
3	31 30 152	5	26 27 332			2	43 46 273			5	11 12 133			5 10 11 133	3	14	13 196	
4	9 6 301	6	17 18 95			3	10 8 267			6				6 11 12 133	4	7	10 87	
5	17 18 47		9,6,L								10,0,L			10,4,L	5			
6	27 28 233	0	53 50 90			5				0	72 63 270					10,8,L		
7	14 15 247	1	44 40 180			1	106 95 270			2	18 17 180			0 46 44 0	0	7	8 180	
	9,3,L	2	85 84 178			3	49 45 270			1	10 8			1 78 77 327	1	30	29 287	
0	37 36 90	0	29 30 270			6				2	15 14 103			2 22 22 52	2	9	6 180	
1	14 14 203	1	18 19 141			4	12 15 103			3	12 15 103			3 15 14 68	3	6	4 311	
2	100 100 124	2	27 29 67			5	16 17 154			4	8 9 261			4 12 15 103	4	17	18 317	
3	61 61 97	3	16 14 98			0	11 3 0			5	10,1,L			5 16 16 344	5	11	12 290	
4	40 42 170	4	15 16 165			1	75 72 326			0	11 75 326			10,5,L		10,9,L		
5	16 16 230	5	24 25 262			2	21 21 17			1	21 21 17				0	23	24 180	
6	11 11 331	6	15 16 165			3	35 36 20			2	12 11 32			10,5,L	0	10	7 302	
7	17 21 51	0	7 2 90			4	24 22 90			3	9 9 185				1	18	19 305	
	9,4,L	1	39 41 194			5	24 22 90			4	24 26 351			10,5,L	2	5	5 118	
0	110 114 90	2	73 76 88			0	24 22 90			0	10,2,L				3			
1	42 40 260	3	47 45 304			1	19 21 341			1	65 63 180			10,6,L	0	8	11 0	
2	50 50 130	4	10 8 335			2	21 20 298			2	45 43 15				1	24	24 211	
3	36 36 106	5	16 15 29			3	5 3 357			3	25 25 275				2	24	24 183	
4	42 44 198	6	13 12 283			4	6 7 113			4	21 24 66				3	23	24 280	
5	20 20 209	0	9,8,L			0	9,12,L			5	13 15 36							
		0	7 6 90			1	12 13 90				13 15 36							

0	10,11,L	11,2,L	4	6	6	130	11,10,L	12,3,L	12,8,L
1	15 15 0	20 20 90		11,6,L			0 19 18 270	0 14 13 180	1 10 11 342
2	12 13 345	7 9 235	0	20 22 90	2	15 14 118	1 6 5 209	1 14 14 343	
3	9 7 351	18 17 4	1	11 12 226	2	15 14 118	2 6 5 332	2 8 8 233	13,0,L
	10,12,L	18 18 233	2	14 14 267		12,0,L	4 13 10 241	1 8 7 90	
0	25 25 180	11,3,L	3	5 1 214	0	15 17 0	12,4,L	2 38 30 180	
1	13 14 212	17 18 90	4	22 22 115	1	17 18 270			13,1,L
	11,0,L	12 11 210		11,7,L	2	21 21 180	0 9 10 180	0 9 10 90	
2	45 43 90	42 44 102	0	18 19 90	3	10 10 270	1 27 28 341	1 16 13 36	
3	7 0 90	11 9 102	1	6 6 318	4	10 10 0	2 10 11 230	2 22 17 104	
4	13 14 180	11,4,L	2	16 16 85		12,1,L	3 16 16 230		13,2,L
5	8 10 270	12 6 90	4	15 15 327	0	11 10 180			
	11,1,L	20 18 24		11,8,L	2	10 9 289	2 17 16 229	0 12 10 90	
0	55 58 90	30 31 187	0	6 5 270	3	22 21 11	3 12 10 115	1 19 17 153	
1	18 17 179	24 25 240	1	6 7 186	4	7 8 100			13,3,L
2	18 16 122	18 17 126	2	10 7 186		12,2,L	12,6,L		
3	29 30 134		3	29 28 93	0	23 25 180	0 20 20 0	0 11 11 90	
4	10 8 131	11,5,L		11,9,L	1	25 27 241	2 18 16 271		13,4,L
5	13 12 201	35 34 90	0	9 9 270	2	26 26 16	3 9 8 293		
		7 5 57	1	16 16 60	3	10 10 25	0 12,7,L	0 7 7 270	
		17 19 47	2	9 9 37	4	9 7 44	1 13 13 180	1 8 7 111	
		7 134	3	12 12 191					

Table XXIII Mikanokryptin: Final structure factors

The table shows:

L^*	$ F_o $	$ F_c $	Phase($^\circ$)
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* Reflections sorted into groups with common
H,K indices (shown in each group heading).

[illegible]

0	1	2	3	4	5	6	7	8	9	-1,-9,L	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108	109	110	111	112	113	114	115	116	117	118	119	120	121	122	123	124	125	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160	161	162	163	164	165	166	167	168	169	170	171	172	173	174	175	176	177	178	179	180	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	206	207	208	209	210	211	212	213	214	215	216	217	218	219	220	221	222	223	224	225	226	227	228	229	230	231	232	233	234	235	236	237	238	239	240	241	242	243	244	245	246	247	248	249	250	251	252	253	254	255	256	257	258	259	260	261	262	263	264	265	266	267	268	269	270	271	272	273	274	275	276	277	278	279	280	281	282	283	284	285	286	287	288	289	290	291	292	293	294	295	296	297	298	299	300	301	302	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338	339	340	341	342	343	344	345	346	347	348	349	350	351	352	353	354	355	356	357	358	359	360	361	362	363	364	365	366	367	368	369	370	371	372	373	374	375	376	377	378	379	380	381	382	383	384	385	386	387	388	389	390	391	392	393	394	395	396	397	398	399	400	401	402	403	404	405	406	407	408	409	410	411	412	413	414	415	416	417	418	419	420	421	422	423	424	425	426	427	428	429	430	431	432	433	434	435	436	437	438	439	440	441	442	443	444	445	446	447	448	449	450	451	452	453	454	455	456	457	458	459	460	461	462	463	464	465	466	467	468	469	470	471	472	473	474	475	476	477	478	479	480	481	482	483	484	485	486	487	488	489	490	491	492	493	494	495	496	497	498	499	500	501	502	503	504	505	506	507	508	509	510	511	512	513	514	515	516	517	518	519	520	521	522	523	524	525	526	527	528	529	530	531	532	533	534	535	536	537	538	539	540	541	542	543	544	545	546	547	548	549	550	551	552	553	554	555	556	557	558	559	560	561	562	563	564	565	566	567	568	569	570	571	572	573	574	575	576	577	578	579	580	581	582	583	584	585	586	587	588	589	590	591	592	593	594	595	596	597	598	599	600	601	602	603	604	605	606	607	608	609	610	611	612	613	614	615	616	617	618	619	620	621	622	623	624	625	626	627	628	629	630	631	632	633	634	635	636	637	638	639	640	641	642	643	644	645	646	647	648	649	650	651	652	653	654	655	656	657	658	659	660	661	662	663	664	665	666	667	668	669	670	671	672	673	674	675	676	677	678	679	680	681	682	683	684	685	686	687	688	689	690	691	692	693	694	695	696	697	698	699	700	701	702	703	704	705	706	707	708	709	710	711	712	713	714	715	716	717	718	719	720	721	722	723	724	725	726	727	728	729	730	731	732	733	734	735	736	737	738	739	740	741	742	743	744	745	746	747	748	749	750	751	752	753	754	755	756	757	758	759	760	761	762	763	764	765	766	767	768	769	770	771	772	773	774	775	776	777	778	779	780	781	782	783	784	785	786	787	788	789	790	791	792	793	794	795	796	797	798	799	800	801	802	803	804	805	806	807	808	809	810	811	812	813	814	815	816	817	818	819	820	821	822	823	824	825	826	827	828	829	830	831	832	833	834	835	836	837	838	839	840	841	842	843	844	845	846	847	848	849	850	851	852	853	854	855	856	857	858	859	860	861	862	863	864	865	866	867	868	869	870	871	872	873	874	875	876	877	878	879	880	881	882	883	884	885	886	887	888	889	890	891	892	893	894	895	896	897	898	899	900	901	902	903	904	905	906	907	908	909	910	911	912	913	914	915	916	917	918	919	920	921	922	923	924	925	926	927	928	929	930	931	932	933	934	935	936	937	938	939	940	941	942	943	944	945	946	947	948	949	950	951	952	953	954	955	956	957	958	959	960	961	962	963	964	965	966	967	968	969	970	971	972	973	974	975	976	977	978	979	980	981	982	983	984	985	986	987	988	989	990	991	992	993	994	995	996	997	998	999	1000	1001	1002	1003	1004	1005	1006	1007	1008	1009	1010	1011	1012	1013	1014	1015	1016	1017	1018	1019	1020	1021	1022	1023	1024	1025	1026	1027	1028	1029	1030	1031	1032	1033	1034	1035	1036	1037	1038	1039	1040	1041	1042	1043	1044	1045	1046	1047	1048	1049	1050	1051	1052	1053	1054	1055	1056	1057	1058	1059	1060	1061	1062	1063	1064	1065	1066	1067	1068	1069	1070	1071	1072	1073	1074	1075	1076	1077	1078	1079	1080	1081	1082	1083	1084	1085	1086	1087	1088	1089	1090	1091	1092	1093	1094	1095	1096	1097	1098	1099	1100	1101	1102	1103	1104	1105	1106	1107	1108	1109	1110	1111	1112	1113	1114	1115	1116	1117	1118	1119	1120	1121	1122	1123	1124	1125	1126	1127	1128	1129	1130	1131	1132	1133	1134	1135	1136	1137	1138	1139	1140	1141	1142	1143	1144	1145	1146	1147	1148	1149	1150	1151	1152	1153	1154	1155	1156	1157	1158	1159	1160	1161	1162	1163	1164	1165	1166	1167	1168	1169	1170	1171	1172	1173	1174	1175	1176	1177	1178	1179	1180	1181	1182	1183	1184	1185	1186	1187	1188	1189	1190	1191	1192	1193	1194	1195	1196	1197	1198	1199	1200	1201	1202	1203	1204	1205	1206	1207	1208	1209	1210	1211	1212	1213	1214	1215	1216	1217	1218	1219	1220	1221	1222	1223	1224	1225	1226	1227	1228	1229	1230	1231	1232	1233	1234	1235	1236	1237	1238	1239	1240	1241	1242	1243	1244	1245	1246	1247	1248	1249	1250	1251	1252	1253	1254	1255	1256	1257	1258	1259	1260	1261	1262	1263	1264	1265	1266	1267	1268	1269	1270	1271	1272	1273	1274	1275	1276	1277	1278	1279	1280	1281	1282	1283	1284	1285	1286	1287	1288	1289	1290	1291	1292	1293	1294	1295	1296	1297	1298	1299	1300	1301	1302	1303	1304	1305	1306	1307	1308	1309	1310	1311	1312	1313	1314	1315	1316	1317	1318	1319	1320	1321	1322	1323	1324	1325	1326	1327	1328	1329	1330	1331	1332	1333	1334	1335	1336	1337	1338	1339	1340	1341	1342	1343	1344	1345	1346	1347	1348	1349	1350	1351	1352	1353	1354	1355	1356	1357	1358	1359	1360	1361	1362	1363	1364	1365	1366	1367	1368	1369	1370	1371	1372	1373	1374	1375	1376	1377	1378	1379	1380	1381	1382	1383	1384	1385	1386	1387	1388	1389	1390	1391	1392	1393	1394	1395	1396	1397	1398	1399	1400	1401	1402	1403	1404	1405	1406	1407	1408	1409	1410	1411	1412	1413	1414	1415	1416	1417	1418	1419	1420	1421	1422	1423	1424	1425	1426	1427	1428	1429	1430	1431	1432	1433	1434	1435	1436	1437	1438	1439	1440	1441	1442	1443	1444	1445	1446	1447	1448	1449	1450	1451	1452	1453	1454	1455	1456	1457	1458	1459	1460	1461	1462	1463	1464	1465	1466	1467	1468	1469	1470	1471	1472	1473	1474	1475	1476	1477	1478	1479	1480	1481	1482	1483	1484	1485	1486	1487	1488	1489	1490
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11	66	69	197							9	60	61	103							2	48	48	270							2	40	40	246							11	171	183	148							11	62	65	16																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																				
10	117	124	77							8	63	60	16							1	97	103	215							1	32	30	96							10	94	95	66							10	83	88	1																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																				
9	143	151	316							7	82	80	297							0	119	124	271							0	28	24	71							9	28	24	63							9	177	188	169																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																				
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-4,-2,L										-4,-4,L										-4,-6,L										-4,-7,L										-4,-9,L										-4,-10,L										-4,-11,L										-4,-12,L									
17	30	27	311	6	150	151	249	13	25	27	37	16	31	30	174	14	23	29	144	11	41	42	202	8	15	11	159	11	41	42	202																																																
16	36	31	235	5	123	120	152	12	52	55	214	15	56	58	148	13	23	19	20	10	22	22	41	7	26	19	123	10	22	22	41																																																
15	39	43	359	4	109	100	222	11	45	48	206	14	39	37	141	12	23	19	20	9	43	40	165	6	26	19	123	9	43	40	165																																																
14	103	105	6	3	161	156	153	10	63	60	18	13	57	58	17	11	68	66	349	8	18	18	167	5	31	37	342	8	18	18	167																																																
13	133	136	144	2	233	226	256	9	80	83	156	10	49	50	148	12	68	66	349	7	17	12	145	4	22	25	254	7	17	12	145																																																
12	69	68	263	1	32	35	12	8	184	194	228	11	49	50	148	11	30	28	198	6	20	19	163	3	31	37	342	6	20	19	163																																																
11	98	103	113	0	21	15	0	7	40	42	3	9	100	105	53	9	31	33	244	5	45	44	225	2	34	30	61	5	45	44	225																																																
10	116	115	262	-4,-4,L						6	32	32	23	10	60	57	292	11	30	28	198	4	71	68	109	4	71	68	109																																																		
9	139	139	130	18	55	55	88	5	45	44	355	9	42	44	206	8	34	36	121	3	27	29	110	3	27	29	110	3	27	29	110																																																
8	154	162	316	17	24	24	92	4	145	145	73	7	118	122	268	6	97	101	232	2	34	30	61	2	34	30	61	2	34	30	61																																																
7	38	41	148	15	63	59	227	3	143	150	226	6	97	100	241	5	15	19	51	1	24	27	76	1	24	27	76	1	24	27	76																																																
6	175	171	230	14	64	64	342	2	169	170	166	5	111	111	352	4	78	80	50	0	67	67	180	0	67	67	180	0	67	67	180																																																
5	141	135	142	13	52	50	39	1	180	190	0	4	68	70	19	3	90	90	41	2	67	67	180	0	67	67	180	0	67	67	180																																																
4	136	130	25	12	87	86	161	0	-4,-6,L						2	65	72	311	4	90	90	41	1	67	67	180	0	67	67	180	0	67	67	180																																													
3	204	194	134	11	111	114	60	1	-4,-6,L						3	204	217	291	3	37	37	270	2	67	67	180	0	67	67	180	0	67	67	180																																													
2	258	242	273	10	54	50	291	2	-4,-6,L						2	204	217	291	2	37	37	270	1	67	67	180	0	67	67	180	0	67	67	180																																													
1	442	419	32	9	82	83	201	16	56	53	88	1	70	63	196	1	57	60	228	0	-4,-12,L						0	67	67	180	0	67	67	180																																													
0	100	98	180	8	141	144	303	15	29	32	340	0	105	113	0	0	106	108	180	-4,-10,L						8	15	11	159	8	15	11	159																																														
-4,-3,L										7	97	95	144	13	15	19	262	1	105	113	0	-4,-10,L						7	74	71	69	7	74	71	69																																												
18	35	38	271	6	199	195	254	12	33	38	183	14	64	62	15	12	60	60	187	6	26	19	123	6	26	19	123	6	26	19	123																																																
17	88	89	211	5	205	202	183	11	62	61	127	13	24	22	266	11	24	28	357	5	31	30	13	5	31	30	13	5	31	30	13																																																
16	63	62	107	4	53	49	63	10	40	40	64	14	64	62	15	10	24	28	357	4	39	37	342	4	39	37	342	4	39	37	342																																																
15	17	18	175	3	401	406	114	9	25	21	191	13	24	22	266	10	24	28	357	3	31	30	13	3	31	30	13	3	31	30	13																																																
14	23	23	308	2	17	17	117	8	45	48	299	11	58	55	2	9	32	37	62	2	26	25	56	2	26	25	56	2	26	25	56																																																
13	93	91	172	1	73	80	127	7	111	112	349	10	60	56	65	8	32	37	62	1	20	15	159	5	20	15	159	5	20	15	159																																																
12	79	78	355	0	67	71	0	6	119	121	117	9	61	56	65	7	51	51	73	0	-4,-13,L						5	20	15	159	5	20	15	159																																													
11	51	50	345	-4,-5,L						5	45	38	337	8	60	56	65	6	58	59	270	-4,-13,L						5	20	15	159	5	20	15	159																																												
10	91	100	134	17	43	43	223	4	200	202	96	7	38	36	325	6	39	40	61	3	30	28	80	3	30	28	80	3	30	28	80																																																
9	161	169	194	16	63	61	186	3	122	124	85	6	43	44	230	5	60	61	114	2	38	34	237	2	38	34	237	2	38	34	237																																																
8	97	96	115	15	62	62	120	2	50	53	14	5	54	55	60	4	88	87	106	1	33	26	27	1	33	26	27	1	33	26	27																																																
7	45	47	75	14	56	59	321	1	93	98	180	4	24	22	119	3	24	22	119	0	20	24	180	0	20	24	180	0	20	24	180																																																

-5,0,L									
17	15	8	90	3	294	293	16	10	49
16	45	47	180	2	157	145	176	9	124
15	44	47	90	1	172	164	284	8	128
13	18	7	90	0	53	53	90	7	104
12	47	50	0	-5,-2,L					
11	100	108	90	17	35	36	246	4	267
10	69	78	180	16	24	26	126	3	379
9	269	285	90	15	59	58	81	2	294
8	21	24	0	14	34	37	70	1	122
7	53	56	90	13	59	59	312	0	230
6	100	102	0	12	16	11	99	-5,-4,L	
5	93	93	90	11	126	131	69	16	59
4	251	244	180	10	143	143	203	15	33
3	251	244	90	9	126	129	99	14	84
2	57	58	180	8	99	97	72	13	33
1	229	207	90	7	150	157	124	12	56
-5,-1,L									
18	55	53	283	6	73	71	298	11	45
17	30	26	342	5	162	165	70	10	78
16	47	46	337	4	309	313	227	9	75
14	28	24	67	3	155	146	228	8	32
13	16	14	326	2	22	21	174	7	31
12	36	34	52	1	237	236	152	6	66
11	14	23	249	0	226	224	90	5	45
10	137	150	164	-5,-3,L					
9	222	226	327	18	17	6	144	4	148
8	65	68	41	16	47	46	280	3	154
7	199	200	330	15	21	21	341	2	164
6	73	75	235	14	36	37	99	1	238
5	76	75	319	13	44	44	339	0	92
4	46	48	91	12	34	34	91	-5,-5,L	
17	52	53	74	-5,-6,L					
16	22	14	304	16	59	59	150	16	59
15	25	21	293	15	24	23	153	15	24
14	27	21	185	14	29	32	48	14	29
13	29	31	72	13	15	18	218	13	15
12	44	44	203	12	71	70	8	12	71
11	37	38	239	11	46	47	10	11	46
10	69	74	50	10	51	54	137	10	51
9	110	112	34	9	46	47	335	9	46
8	58	61	119	8	116	119	241	8	116
7	135	145	340	7	59	66	342	7	59
6	80	86	211	6	117	121	348	6	117
5	36	31	228	5	106	113	305	5	106
4	160	164	165	4	27	26	24	4	27
3	70	70	119	3	142	147	127	3	142
2	56	56	73	2	47	47	310	2	47
1	66	70	270	1	47	47	310	1	47
-5,-7,L									
16	30	27	90	16	59	59	150	16	59
15	25	25	260	15	24	23	153	15	24
14	52	49	50	14	29	32	48	14	29
13	26	27	57	13	15	18	218	13	15
12	37	35	198	12	71	70	8	12	71
11	27	28	236	11	46	47	10	11	46
10	51	49	140	10	51	54	137	10	51
9	39	43	359	9	46	47	335	9	46
8	19	17	1	8	116	119	241	8	116
7	90	94	291	7	59	66	342	7	59
6	49	50	129	6	117	121	348	6	117
5	19	17	1	5	106	113	305	5	106
4	49	50	49	4	27	26	24	4	27
3	173	180	49	3	142	147	127	3	142
2	68	67	136	2	47	47	310	2	47
1	106	113	5	1	47	47	310	1	47
-5,-8,L									
15	52	50	122	16	59	59	150	16	59
14	42	42	26	15	24	23	153	15	24
13	52	53	36	14	29	32	48	14	29
12	43	45	171	13	15	18	218	13	15
11	27	28	143	12	71	70	8	12	71
10	38	40	29	11	46	47	10	11	46
9	62	61	189	10	51	54	137	10	51
8	50	50	294	9	46	47	335	9	46
7	77	80	69	8	116	119	241	8	116
6	45	46	349	7	59	66	342	7	59
5	88	94	318	6	117	121	348	6	117
4	22	22	234	5	106	113	305	5	106
3	88	94	318	4	27	26	24	4	27
2	22	22	234	3	142	147	127	3	142
1	47	47	310	2	47	47	310	2	47
-5,-9,L									
13	15	6	258	16	59	59	150	16	59
12	54	51	19	15	24	23	153	15	24
10	52	55	336	14	29	32	48	14	29
9	32	32	88	13	15	18	218	13	15
8	29	32	9	12	71	70	8	12	71
7	35	37	84	11	46	47	10	11	46
6	66	67	358	10	51	54	137	10	51
5	39	40	316	9	46	47	335	9	46
4	22	20	111	8	116	119	241	8	116
3	104	101	30	7	59	66	342	7	59
2	76	76	73	6	117	121	348	6	117
1	112	117	270	5	106	113	305	5	106
-5,-10,L									
11	25	26	28	16	59	59	150	16	59
10	19	20	158	15	24	23	153	15	24
9	14	18	290	14	29	32	48	14	29
8	18	13	298	13	15	18	218	13	15
7	46	46	182	12	71	70	8	12	71
6	44	44	229	11	46	47	10	11	46
5	34	34	90	10	51	54	137	10	51
4	42	43	94	9	46	47	335	9	46
3	25	25	321	8	116	119	241	8	116
2	15	9	270	7	59	66	342	7	59
1	42	43	94	6	117	121	348	6	117
-5,-11,L									
10	24	15	283	16	59	59	150	16	59
9	28	25	21	15	24	23	153	15	24
8	53	52	62	14	29	32	48	14	29

[illegible]

-6,-9,L									
8	35	34	166						
6	14	16	126						
5	73	74	262						
4	45	46	24						
3	54	53	89						
2	31	29	56						
1	24	24	202						
0	42	39	0						
-6,-10,L									
11	62	62	328						
10	19	19	126						
9	23	25	0						
7	22	22	34						
6	32	28	239						
5	40	38	353						
4	23	20	339						
3	23	27	79						
2	64	63	23						
1	41	41	334						
0	58	62	0						
-6,-11,L									
8	40	38	224						
6	29	33	65						
5	37	35	219						
4	33	30	321						
3	29	30	174						
2	76	74	82						
1	16	21	118						
-6,-12,L									
7	36	28	359						
5	15	17	325						
2	26	26	110						
-6,-13,L									
1	38	36	63						
-7,0,L									
17	46	46	270						
16	38	35	180						
15	82	78	270						
13	84	86	90						
12	50	52	0						
11	66	63	270						
10	97	97	180						
9	200	212	90						
8	15	11	0						
7	19	21	270						
6	72	73	0						
5	91	93	270						
2	97	102	180						
1	31	37	270						
-7,-1,L									
16	64	62	293						
15	36	38	145						
14	49	50	265						
13	47	47	184						
12	54	57	265						
11	20	23	18						
-7,-2,L									
17	16	12	302						
15	34	34	280						
14	40	39	326						
13	20	21	263						
12	56	53	80						
11	107	109	323						
10	52	54	273						
9	92	95	96						
8	111	114	103						
7	89	89	66						
6	127	134	333						
5	126	127	283						
4	62	60	238						
3	96	103	75						
2	67	66	331						
1	151	150	162						
0	80	85	90						
-7,-3,L									
17	28	25	197						
-7,-4,L									
16	24	25	298						
15	65	68	293						
14	94	94	61						
13	33	33	6						
12	90	92	53						
11	17	14	275						
10	86	84	331						
9	27	29	314						
8	69	72	334						
7	73	76	170						
6	115	120	21						
5	157	167	262						
4	75	79	287						
3	102	101	146						
2	68	70	351						
1	62	64	282						
-7,-5,L									
16	28	28	158						
15	30	27	43						
14	28	27	259						
13	31	28	225						
12	51	50	152						
11	19	15	295						
10	31	32	28						
9	40	42	110						
8	30	29	29						
7	41	41	214						
6	88	98	203						
5	40	41	258						
4	106	106	274						
3	145	149	177						
2	157	169	248						
1	53	57	67						
0	220	225	270						
-7,-6,L									
15	51	51	267						
14	46	43	322						
13	37	36	87						
12	58	52	14						
11	38	39	323						
9	24	22	320						
8	62	62	34						
7	63	62	292						
6	54	55	357						
5	35	35	266						
4	142	147	275						
-7,-7,L									
14	28	32	91						
13	34	32	354						
11	26	29	160						
10	62	62	346						
9	72	73	189						
8	43	43	354						
7	24	27	181						
6	86	85	229						
5	35	36	0						
4	29	31	226						
2	74	78	217						
1	15	17	102						
-7,-8,L									
13	20	15	252						
12	40	42	101						
11	31	29	306						
10	33	34	210						
9	41	40	224						
8	42	40	167						
7	15	8	192						
6	64	61	266						
5	35	39	30						
4	52	54	182						
3	51	51	359						
2	13	14	69						

-8,-9,L											
7	22	24	177	15	29	27	270	15	41		
6	33	33	3	14	44	44	180	14	39		
5	63	62	226	13	31	29	270	13	58		
4	20	21	307	12	33	38	180	12	28		
3	49	50	139	10	79	79	180	11	41		
2	24	25	37	8	16	14	180	9	18		
1	51	51	96	7	70	78	270	8	87		
0	29	35	0	6	24	22	180	7	37		
-8,-10,L											
9	17	9	15	4	172	177	0	6	47		
8	33	37	226	3	150	156	270	5	93		
7	37	35	205	2	19	20	180	4	69		
6	49	47	162	1	116	123	90	3	38		
5	23	21	111	-9,-1,L						2	54
4	26	24	272	15	30	27	178	1	137		
3	43	43	51	14	65	64	260	0	136		
2	26	28	0	13	27	24	241	-9,-3,L			
1	43	43	51	12	21	17	177	15	32		
0	26	28	0	11	43	43	358	14	40		
-8,-11,L											
4	28	31	305	9	39	38	168	13	37		
3	27	23	200	8	16	21	78	12	40		
2	48	47	77	7	21	26	202	11	51		
1	29	32	143	6	86	92	112	10	40		
0	29	31	0	5	75	77	261	9	36		
-8,-12,L											
1	23	23	134	4	124	129	128	8	41		

-10,0,L									
11	35	35	270	13	41	41	284	12	47
10	34	37	0	12	46	44	314	11	56
9	28	24	270	11	19	19	179	10	15
7	26	30	90	10	19	5	251	9	25
5	18	12	270	9	38	35	37	8	18
4	61	60	0	8	35	36	268	7	15
3	36	36	90	7	46	46	53	5	109
2	90	96	0	6	39	36	28	4	77
1	97	97	90	5	110	109	70	3	70
0	55	55	0	4	136	139	310	2	49
-10,-1,L									
15	18	19	17	3	14	16	121	1	74
14	48	47	324	2	68	72	353		
13	63	61	217	1	23	26	286		
12	25	21	113	0	25	27	180		
11	33	28	18	-10,-3,L					
10	57	54	39	13	25	28	229	13	20
9	75	72	179	12	24	24	246	12	16
8	95	97	195	11	40	42	205	11	31
7	43	44	152	10	66	64	35	10	59
6	61	63	214	9	46	49	352	9	89
5	78	77	350	8	18	20	357	8	50
4	27	28	328	7	80	77	198	7	38
3	30	15	7	6	31	35	112	6	28
2	39	44	124	5	34	35	90	5	50
1	16	19	74	4	34	33	223	4	91
0	43	39	180	3	31	31	350	3	50
-10,-2,L									
14	30	29	256	2	86	93	181	2	34
				1	37	41	107	1	59
				0	151	151	180		
-10,-4,L									
13	19	20	178	-10,-6,L					
				12	23	24	89	12	23
				10	75	72	46	10	75
				8	77	72	91	8	77
				7	49	49	26	7	49
-10,-7,L									
				6	21	23	21	6	21
				4	51	51	45	4	80
				3	80	80	270	3	41
				2	41	39	300	2	45
				1	45	44	78	1	17
				0	17	14	180		
-10,-8,L									
				11	26	32	356	11	26
				10	46	46	150	10	21
				9	21	17	94	9	31
				8	31	35	83	8	33
				7	33	31	37	7	19
				6	19	21	16	6	19
				5	19	23	246	5	44
				4	44	40	48	4	22
				3	22	30	194	3	66
				2	66	66	9	2	30
				1	30	32	39	1	15
				0	15	15	0		
-10,-9,L									
				10	43	42	26	10	43
				7	32	30	128	7	58
				6	58	55	70	6	35
				4	35	34	356	4	33
				3	33	31	269	3	43
				2	43	46	259	2	57
				1	57	53	306	1	14
				0	14	15	180		
-11,0,L									
13	17	11	270	13	17	11	270	13	17
12	21	21	180	12	21	21	180	12	21
11	47	39	90	11	47	39	90	11	47
10	30	27	180	10	30	27	180	10	30
9	44	44	270	9	44	44	270	9	44
8	81	81	0	8	81	81	0	8	81
7	45	44	90	7	45	44	90	7	45
6	41	39	0	6	41	39	0	6	41
5	48	52	90	5	48	52	90	5	48
4	81	81	0	4	81	81	0	4	81
3	45	44	90	3	45	44	90	3	45
2	41	39	0	2	41	39	0	2	41
1	48	52	90	1	48	52	90	1	48
0	81	81	0	0	81	81	0	0	81
-11,-1,L									
12	46	45	274	12	46	45	274	12	46
11	29	27	230	11	29	27	230	11	29
10	15	15	42	10	15	15	42	10	15
9	22	16	256	9	22	16	256	9	22
8	32	30	309	8	32	30	309	8	32
7				7				7	
-11,-3,L									
13	38	35	358	13	38	35	358	13	38
12	51	48	312	12	51	48	312	12	51
11	49	44	269	11	49	44	269	11	49
10	53	51	315	10	53	51	315	10	53
9	31	28	47	9	31	28	47	9	31
8	66	64	200	8	66	64	200	8	66
7	52	52	18	7	52	52	18	7	52
6	40	40	310	6	40	40	310	6	40
5	50	48	328	5	50	48	328	5	50
4	73	72	358	4	73	72	358	4	73
3				3				3	

-13,-2,L

8 21 21 197
7 35 33 263
6 24 26 195
5 31 30 8
4 41 41 192
3 30 26 265
2 58 54 23
1 26 27 281
0 24 23 270

-13,-3,L

9 34 31 178
7 38 36 205
6 39 36 292
5 33 33 194
4 36 38 354
3 33 31 258
2 33 21 241
1 21 25 90
0 28

-13,-4,L

8 31 28 16
7 22 22 244
6 27 23 9
4 35 31 90
2 59 56 2
1 48 48 34
0 74 75 270

-13,-5,L

8 28 23 102
5 43 40 240
4 37 34 321
2 40 42 289
1 48 47 234
0 21 24 270

-13,-6,L

6 38 37 340
4 15 7 128
3 41 41 154

2 22 23 9
1 42 39 91

-13,-7,L

4 27 17 343
2 33 32 78
1 30 34 225
0 17 19 90

-14,0,L

8 16 13 180
7 28 27 90
5 25 26 90
4 48 47 180
2 92 86 180
1 36 35 90
0 73 71

-14,-1,L

8 36 36 278
7 37 35 195

6 35 33 314
5 72 69 176
2 25 21 310
1 20 18 51
0 17 15 180

-14,-2,L

7 18 15 128
4 16 19 125
3 20 22 345
2 33 27 187
1 62 58 146
0 36 33

-14,-3,L

6 18 19 254
5 20 17 54
4 48 44 116
3 26 25 96
2 21 18 315
1 57 58 328
0 28 25

-14,-4,L

6 16 20 296
5 23 22 338
4 27 27 268
3 27 23 17
2 45 46 301
0 19 14 0

-14,-5,L

4 22 19 95
3 28 25 117
2 27 22 101
1 39 39 297
0 46 46 0

-15,0,L

4 29 27 180
3 21 20 90
2 18 12 180
1 31 28 270

-15,-1,L

4 40 37 266
3 39 38 197
2 28 26 141
1 22 18 356

-15,-2,L

3 23 19 149
1 15 15 222
0 45 43 270

-15,-3,L

2 35 32 98
1 24 22 344
0 25 25 90

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CHAPTER THREE

An x-ray analysis of two derivatives of 2-naphthol

3.1 Introduction

Substituents in the 1,8-or 4, 5- positions in naphthalene (I) cause greater steric hindrance than if they were ortho (e.g. 1,2-). The interactions resulting from this type of substitution are known as *peri*-interactions and have been the subject of a comprehensive review (1).

The effect of *peri*-strain on the autoxidation of 1-alkyl-2- naphthols has recently been reported by Brady and Carnduff (2). This reaction shows the characteristics of a radical chain process and occurs in the presence of atmospheric oxygen.

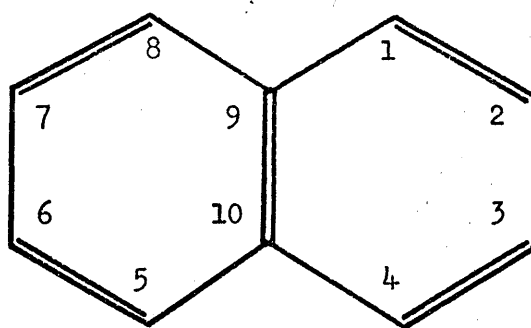
Brady and Carnduff found that the relative times for the uptake of a given quantity of oxygen were as follows:

$$A (\text{inert}) = B \gg C \gg D \gg E \text{ (Fig. 3.1).}$$

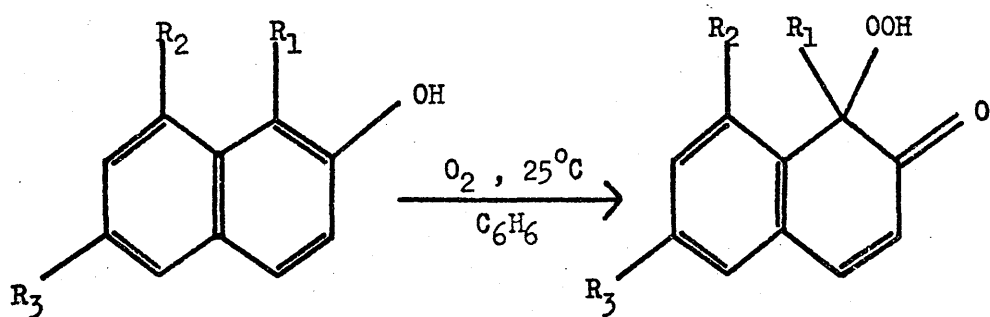
These reaction rates correlate with the increasing strain in the *peri*-positions and may possibly be accounted for by the relief in steric hindrance when C(1) becomes tetrahedral.

Dr. J. Carnduff (University of Glasgow) supplied two of a series of derivatives of 2-naphthols for the following x-ray analyses. These analyses form part of a study to assess the relative strain in a number of compounds in which R_1 , R_2 , R_3 (Fig. 3.1) are allowed to vary.

The crystal structure analyses could not easily be carried out on the 2-naphthols themselves since the autoxidation occurs at room temperature in the presence of molecular oxygen. The 2-hydroxyl group was therefore



(I)



	R_1	R_2	R_3
<u>A</u>	Me	H	H
<u>B</u>	Me	H	Bu ^t
<u>C</u>	Pr ⁱ	H	H
<u>D</u>	Me	Me	H
<u>E</u>	Bu ^t	H	H

Fig. 3.1 Autoxidation of 1-alkyl-2-naphthols

protected by acetylation; it being assumed that this substitution would have a minimal effect on strain elsewhere in the molecule.

3.2 1—Methyl—2—naphthyl—acetate

Experimental

Crystal Data

1—Methyl—2—naphthyl—acetate: $C_{13}H_{12}O_2$

Unit cell dimensions $\underline{a} = 6.455 \text{ \AA}$ $\underline{\alpha} = 90.0^\circ$
 $\underline{b} = 8.325 \text{ \AA}$ $\underline{\beta} = 91.95^\circ$
 $\underline{c} = 20.252 \text{ \AA}$ $\underline{\gamma} = 90.0^\circ$

Space group $P2_1/C (C_2^5h)$

$M = 200 \text{ a.m.u.}$

$V = 1088 \text{ \AA}^3$

$D_o = 1.41 \text{ gmcm}^{-3}$

$D_c = 1.222 \text{ gmcm}^{-3}$

$\mu = 0.88 \text{ cm}^{-1} (\text{Mo-K}\alpha)$

$F(000) = 424$

$Z = 4$

The space group was uniquely determined by the systematic absences:

$h0l$ when $l = 2n$, $0k0$ when $k = 2n$.

The crystal used in this analysis was a small cube of side 1mm.

Data Collection

Precession photographs were used to measure the cell dimensions and to determine the space group. The crystal was then mounted and aligned on the goniometer head of the Hilger diffractometer as described previously. The cell dimensions quoted in the crystal data are those computed during this diffractometer setting up procedure.

The $\theta, 2\theta$ scanning technique was used to obtain intensity data from zirconium filtered Mo-K α radiation. The scan parameters were as follows:

Increment in <u>θ</u>	0.02°
Number of scan steps	40
Time per step	1 sec.

The background count was measured for 20 seconds immediately before and after each peak count. The intensities of two strong reflexions were checked after every 40 intensity measurements and no significant variation in their values was recorded.

These standards were used to place the measured intensities for the hkl and $h\bar{k}\bar{l}$ octants ($2\theta \leq 52^\circ$) on a common scale.

The integrated intensities (I) were reduced to 1682 independent structure amplitudes by exclusion of the data for which $I \leq 2\sigma I$.

Reduced beam intensity was employed to remeasure the reflexions for which $2\theta \leq 12^\circ$ in order to correct for counter saturation errors.

No absorption correction was applied.

Structure Analysis

The structure was elucidated by the direct phase determining program MULTAN. Triplet relationships were derived for the largest 200 of the normalised structure factors ($|E| \geq 1.45$).

The starting set of 7 reflexions (3 origin defining, Table I) was selected by the program on the basis that phases with a probability in excess of 0.95^s were acceptable. Sixteen phase sets were then calculated for the 200 reflexions with the largest value of $|E|$ and an E -map computed from the set showing the highest combined figure of merit revealed all the non-hydrogen atoms.

A structure factor calculation using these co-ordinates gave an R of 45.8%. The carbon and oxygen atom locations were then refined with a full matrix least squares procedure, using the 1508 independant structure amplitudes for which $|F| \geq 7\sigma(|F|)$, and the calculations converged at $R = 26.1\%$ (isotropic vibration) and $R = 15.6\%$ (anistropic thermal motion).

After making allowance for errors caused by counter saturation R was reduced to 12.2%. Subsequent difference syntheses were used to obtain co-ordinates for all the hydrogen atoms and when these atoms were allowed to assume isotropic vibration R fell to 5.1%

A weighting scheme of the form:

$$W = 1/(A + B F_o + C F_o^2) \text{ with } A=0.1728,$$

$$B=-0.0277 \text{ and } C=0.035$$

was applied. The values of the constants A , B and C

TABLE I SUMMARY OF STRUCTURE DETERMINATION

A. INITIAL SET OF PHASES

REFLECTION	PHASE *	MOD. E
2 5 -13 +	360	4.90
2 7 4 +	360	3.43
1 2 2 +	360	3.21
3 6 -12	360 OK 180	4.14
3 5 6	360 OR 180	3.51
1 4 -15	360 OK 180	3.15
4 2 -9	360 OR 180	2.94

B. STATISTICS

	E1	E2	E3	E4	E5	E6
THEORETICAL FOR CENTRIC	0.798	1.00	0.968	31.73	4.55	0.27
THEORETICAL FOR ACENTRIC	0.886	1.00	0.736	36.79	1.83	0.01
FOUND	0.803	1.00	0.991	27.80	5.00	0.9

* IN DEGREES
+ ORIGIN DEFINING

MOD. E MODULUS OF E FOR REFLECTION CONCERNED

E1 MEAN VALUE FOR MODULUS OF E
E2 MEAN VALUE FOR MODULUS OF E**2
E3 MEAN VALUE FOR MODULUS OF ((E**2)-1)
E4 % OF REFLECTIONS WITH MODULUS OF E > 1.0
E5 % OF REFLECTIONS WITH MODULUS OF E > 2.0
E6 % OF REFLECTIONS WITH MODULUS OF E > 3.0

were chosen so that the deviation from constant $w\lambda^2$ over the range of $|F_0|$ and $\sin\theta$ was minimised.

The refinement then converged at $R = 4.6\%$.

No correction was made for absorption.

3.3 Results

Table II 1-Methyl-2-naphthyl acetate: Fractional atomic
coordinates

Positional estimated standard deviations are
shown in parentheses.

The hydrogen atoms are numbered according to the
atoms to which they are attached.

The table shows:

Atom	x	y	z
------	---	---	---

C(1)	5324(3)	0779(2)	1196(1)
C(2)	4623(3)	1732(2)	0693(1)
C(3)	2786(3)	2624(2)	0705(1)
C(4)	1639(3)	2581(2)	1253(1)
C(5)	1146(4)	1637(3)	2398(1)
C(6)	1824(4)	0766(3)	2933(1)
C(7)	3635(4)	-0127(3)	2903(1)
C(8)	4743(4)	-0150(2)	2346(1)
C(9)	4134(3)	0752(2)	1784(1)
C(10)	2288(3)	1658(2)	1810(1)
C(11)	7228(4)	-0215(3)	1137(1)
C(12)	6982(3)	3031(2)	0006(1)
C(13)	7935(6)	2922(4)	-0653(1)
O(14)	5717(2)	1770(2)	0102(1)
O(15)	7262(2)	4061(2)	0405(1)
H(3)	2445(30)	3273(25)	0338(10)
H(4)	0344(36)	3142(27)	1287(10)
H(5)	-0098(37)	2208(29)	2391(11)
H(6)	1038(35)	0785(27)	3324(12)
H(7)	4176(37)	-0722(29)	3296(13)
H(8)	6064(40)	-0748(31)	2325(12)
H(111)	8170(46)	-0041(35)	1474(15)
H(112)	7885(37)	-0055(28)	0720(13)
H(113)	6816(48)	-1414(38)	1180(14)
H(131)	8219(47)	1878(42)	-0761(14)
H(132)	9153(68)	3526(50)	-0656(19)
H(133)	6932(60)	3298(45)	-0976(19)

Table III 1-Methyl-2-naphthyl acetate: Thermal parameters

Carbon and oxygen values are anisotropic.

Hydrogen values are isotropic.

Estimated errors in the last digit are shown
in parentheses.

All figures are multiplied by 10^3

The anisotropic temperature factor expression
used was of the form described by equation (12)
in chapter one, the coefficients being in terms
of the mean square amplitudes of vibration U_{ij} .

The table shows:

Atom	U_{11} or U	U_{22}	U_{33}	U_{12}	U_{13}	U_{23}
------	-----------------	----------	----------	----------	----------	----------

C(1)	68(1)	50(1)	55(1)	-4(1)	5(1)	-3(1)
C(2)	83(1)	50(1)	49(1)	-8(1)	9(1)	-5(1)
C(3)	97(1)	57(1)	58(1)	6(1)	1(1)	6(1)
C(4)	79(1)	63(1)	72(1)	12(1)	3(1)	0(1)
C(5)	84(1)	60(1)	85(1)	-4(1)	25(1)	-7(1)
C(6)	118(2)	66(1)	63(1)	-17(1)	36(1)	-1(1)
C(7)	116(2)	71(1)	58(1)	-5(1)	13(1)	10(1)
C(8)	87(1)	66(1)	63(1)	0(1)	5(1)	8(1)
C(9)	68(1)	48(1)	54(1)	-9(1)	7(1)	-2(1)
C(10)	73(1)	48(1)	59(1)	-6(1)	7(1)	-3(1)
C(11)	70(1)	86(2)	73(1)	7(1)	11(1)	5(1)
C(12)	79(1)	60(1)	55(1)	2(1)	9(1)	10(1)
C(13)	125(2)	93(2)	72(2)	-9(2)	36(2)	4(1)
O(14)	111(1)	62(1)	51(1)	-14(1)	21(1)	-5(1)
O(15)	102(1)	69(1)	68(1)	-16(1)	12(1)	-5(1)
H(3)	39(5)					
H(4)	52(6)					
H(5)	53(6)					
H(6)	56(6)					
H(7)	63(7)					
H(8)	65(7)					
H(111)	80(9)					
H(112)	58(7)					
H(113)	93(10)					
H(131)	85(10)					
H(132)	115(14)					
H(133)	108(13)					

TABLE IV

1-METHYL-2-NAPHTHYL-ACETATE:
INTRAMOLECULAR BONDED DISTANCES (ANGSTROMS)
STANDARD DEVIATIONS (ANGSTROMS) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	DIST.	ATOM1	ATOM2	DIST.
C1	C2	1.356(2)	C12	O14	1.349(2)
C1	C9	1.439(2)	C12	O15	1.188(2)
C1	C11	1.490(2)	C3	H3	0.939(20)
C2	C3	1.400(2)	C4	H4	0.962(23)
C2	O14	1.409(2)	C5	H5	0.932(24)
C3	C4	1.355(2)	C6	H6	0.954(23)
C4	C10	1.417(2)	C7	H7	0.990(25)
C5	C6	1.363(3)	C8	H8	0.990(26)
C5	C10	1.421(2)	C11	H11	0.909(30)
C6	C7	1.388(3)	C11	H12	0.966(25)
C7	C8	1.357(3)	C11	H13	1.037(31)
C8	C9	1.406(2)	C13	H131	0.916(34)
C9	C10	1.413(2)	C13	H132	0.933(43)
C12	C13	1.491(3)	C13	H133	0.956(38)

TABLE V 1-METHYL-2-NAPHTHYL ACETATE: SOME INTRAMOLECULAR NON-BONDED SEPARATIONS

ATOM1	ATOM2	DIST #	ATOM1	ATOM2	DIST #
C(1).....C(8)		2.494	O(15).....H(112)		3.51
C(11).....O(14)		2.817	H(4).....H(5)		2.39
C(11).....H(8)		2.58	H(7).....H(8)		2.35
O(15).....H(3)		3.18	H(8).....H(111)		2.31

IN ANGSTROMS

1. The structure of 1-methyl-2-naphthyl acetate is shown in Figure 1. The molecule is a derivative of naphthalene, with a methyl group at position 1 and an acetate group at position 2. The structure is shown in a perspective view, with the naphthalene ring system in the foreground and the acetate group extending away from it. The atoms are labeled with numbers in parentheses, indicating their positions in the molecule. The structure is shown in a perspective view, with the naphthalene ring system in the foreground and the acetate group extending away from it. The atoms are labeled with numbers in parentheses, indicating their positions in the molecule.

TABLE VI 1-METHYL-2-NAPHTHYL-ACETATE: INTERMOLECULAR CONTACTS OF LESS
THAN 3.85 ANGSTROMS

ATOM1	ATOM2	DIST #	ATOM1	ATOM2	DIST #
C(11).....C(4)	*	3.680	C(12).....O(15)	***	3.729
O(15).....C(4)	*	3.482	O(15).....O(15)	***	3.652
O(15).....C(3)	*	3.791	C(5).....C(8)	***	3.791
C(1).....O(14)	**	3.427	C(9).....C(7)	***	3.767
C(2).....O(14)	**	3.334	C(10).....C(7)	***	3.785
C(11).....O(14)	**	3.356	C(10).....C(8)	***	3.665
C(3).....C(13)	***	3.740	O(15).....C(6)	***	3.682
C(3).....O(15)	***	3.559	O(15).....C(7)	***	3.560

IN ANGSTROMS

THE ASTERISKS(*) REFER TO THE FOLLOWING TRANSFORMATIONS OF THE ATOMIC
COORDINATES

* 1.0+X, Y, Z
** 1.0-X, -Y, -Z
*** 1.0-X, 1.0-Y, -Z
**** -X,-0.5+Y,-0.5-Z

TABLE VII

1-METHYL-2-NAPHTHYL-ACETATE: VALENCY ANGLES (DEGREES)
STANDARD DEVIATIONS (DEGREES) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	ATOM3	ANGLE	ATOM1	ATOM2	ATOM3	ANGLE
C9	C1	C2	117.2(1)	C11	C1	C2	121.3(1)
C11	C1	C9	121.5(1)	O14	C2	C1	119.1(1)
C3	C2	C1	123.9(1)	O14	C2	C3	116.9(1)
C4	C3	C2	119.2(1)	C10	C4	C3	120.8(1)
C10	C5	C6	120.6(2)	C7	C6	C5	119.9(2)
C8	C7	C6	120.8(2)	C9	C8	C7	121.6(2)
C10	C9	C1	119.8(1)	C8	C9	C1	122.3(1)
C10	C9	C8	117.9(1)	C5	C10	C4	121.7(1)
C9	C10	C4	119.1(1)	C9	C10	C5	119.1(1)
O14	C12	C13	110.7(1)	O15	C12	C13	126.5(2)
O15	C12	O14	122.8(1)	C12	O14	C2	117.6(1)
H3	C3	C2	117.9(12)	H3	C3	C4	122.9(12)
H4	C4	C3	123.1(12)	H4	C4	C10	116.1(12)
H5	C5	C6	122.5(13)	H5	C5	C10	116.8(13)
H6	C6	C5	118.9(13)	H6	C6	C7	121.2(13)
H7	C7	C6	120.5(14)	H7	C7	C8	118.6(14)
H8	C8	C7	121.4(14)	H8	C8	C9	116.9(14)
H11	C11	C1	112.6(18)	H12	C11	C1	112.4(14)
H13	C11	C1	108.3(17)	H12	C11	H11	109.3(23)
H13	C11	H11	104.9(24)	H13	C11	H12	109.2(21)
H131	C13	C12	111.3(18)	H132	C13	C12	110.3(24)
H133	C13	C12	107.5(23)	H132	C13	H131	109.5(32)
H133	C13	H131	106.5(29)	H133	C13	H132	111.8(34)

TABLE_VIII

1-METHYL-2-NAPHTHYL-ACETATE: TORSION ANGLES(DEGREES)
STANDARD DEVIATIONS(DEGREES) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	ATOM3	ATOM4	ANGLE	ATOM1	ATOM2	ATOM3	ATOM4	ANGLE
C9	C1	C2	C3	-2.7(2)	C9	C1	C2	O14	-178.7(1)
C11	C1	C2	C3	176.4(1)	C11	C1	C2	O14	0.4(2)
C2	C1	C9	C8	-177.9(1)	C2	C1	C9	C10	1.9(2)
C11	C1	C9	C8	3.0(2)	C11	C1	C9	C10	-177.2(1)
C1	C2	C3	C4	1.6(3)	O14	C2	C3	C4	177.7(1)
C1	C2	O14	C12	-101.2(1)	C3	C2	O14	C12	82.5(2)
C2	C3	C4	C10	0.5(3)	C3	C4	C10	C5	177.9(1)
C3	C4	C10	C9	-1.2(2)	C10	C5	C6	C7	-0.7(3)
C6	C5	C10	C4	-178.6(2)	C6	C5	C10	C9	0.5(3)
C5	C6	C7	C8	-0.2(3)	C6	C7	C8	C9	1.5(3)
C7	C8	C9	C1	178.1(1)	C7	C8	C9	C10	-1.7(2)
C1	C9	C10	C4	0.0(2)	C1	C9	C10	C5	-179.1(1)
C4	C9	C10	C4	179.8(1)	C8	C9	C10	C5	0.7(2)
C13	C12	O14	C2	-177.3(1)	O15	C12	O14	C2	2.9(2)
C2	C1	C11	H111	126.3(20)	C2	C1	C11	H112	2.4(16)
C2	C1	C11	H113	-118.2(17)	C9	C1	C11	H111	-54.6(20)
C4	C1	C11	H112	-178.5(15)	C9	C1	C11	H113	60.9(17)
C1	C2	C3	H3	178.3(13)	O14	C2	C3	H3	-5.6(14)
C2	C3	C4	H4	-178.2(16)	H3	C3	C4	C10	-176.1(14)
H3	C3	C4	H4	5.3(21)	H4	C4	C10	C5	-3.4(15)

DIHEDRAL ANGLES (CONTINUED)

ATOM 1	ATOM 2	ATOM 3	ATOM 4	ANGLE	ATOM 1	ATOM 2	ATOM 3	ATOM 4	ANGLE
O14	C4	C10	C9	177.5(15)	C10	C5	C6	H6	179.1(15)
O15	C5	C6	C7	176.7(17)	H5	C5	C6	H6	-3.5(23)
O15	C5	C10	C4	3.9(16)	H5	C5	C10	C9	-177.1(16)
C5	C6	C7	H7	177.1(16)	H6	C6	C7	C8	180.0(16)
H6	C6	C7	H7	-2.7(23)	C6	C7	C8	H8	177.6(17)
H7	C7	C8	C9	-175.9(16)	H7	C7	C8	H8	0.2(23)
H8	C8	C9	C1	1.8(16)	H8	C8	C9	C10	-178.0(16)
O14	C12	C13	H131	-36.0(21)	O14	C12	C13	H132	-157.7(27)
O14	C12	C13	H133	80.2(24)	O15	C12	C13	H131	143.7(21)
O15	C12	C13	H132	22.0(27)	O15	C12	C13	H133	-100.0(24)

THE ANGLE 1-2-3-4 IS DEFINED AS POSITIVE IF, WHEN VIEWED ALONG THE 2-3 BOND, ATOM 1 HAS TO BE ROTATED CLOCKWISE TO ECLIPSE ATOM 4

TABLE IX 1-METHYL-2-NAPHTHYL-ACETATE: MEAN PLANES CALCULATIONS

ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
C(1)		-0.012(2)
C(2)		0.031(2)
C(3)		0.013(2)
C(4)		-0.016(2)
C(5)		-0.004(2)
C(6)		0.021(2)
C(7)		0.018(2)
C(8)		-0.016(2)
C(9)		-0.017(2)
C(10)		-0.019(2)
	C(11)	-0.071(3)
	O(14)	0.010(1)
	H(4)	-0.056(23)
	H(5)	-0.051(25)
	H(8)	0.018(26)

1.

TABLE IX (CONTINUED)

	ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
2.	C(12) C(13) C(14)	0(15)	0.000(2) 0.000(4) 0.000(1) 0.004(2)

THE OUT OF PLANE ATOMS WERE NOT INCLUDED IN THE DERIVATION OF THE PLANES.

THE EQUATIONS OF THESE PLANES ARE:

1. $0.5113X + 0.7815Y + 0.3576Z = 3.0991$
2. $0.7539X - 0.5148Y + 0.4082Z = 2.1024$



Fig. 3.2 1-Methyl-2-naphthyl acetate: A view of the molecule



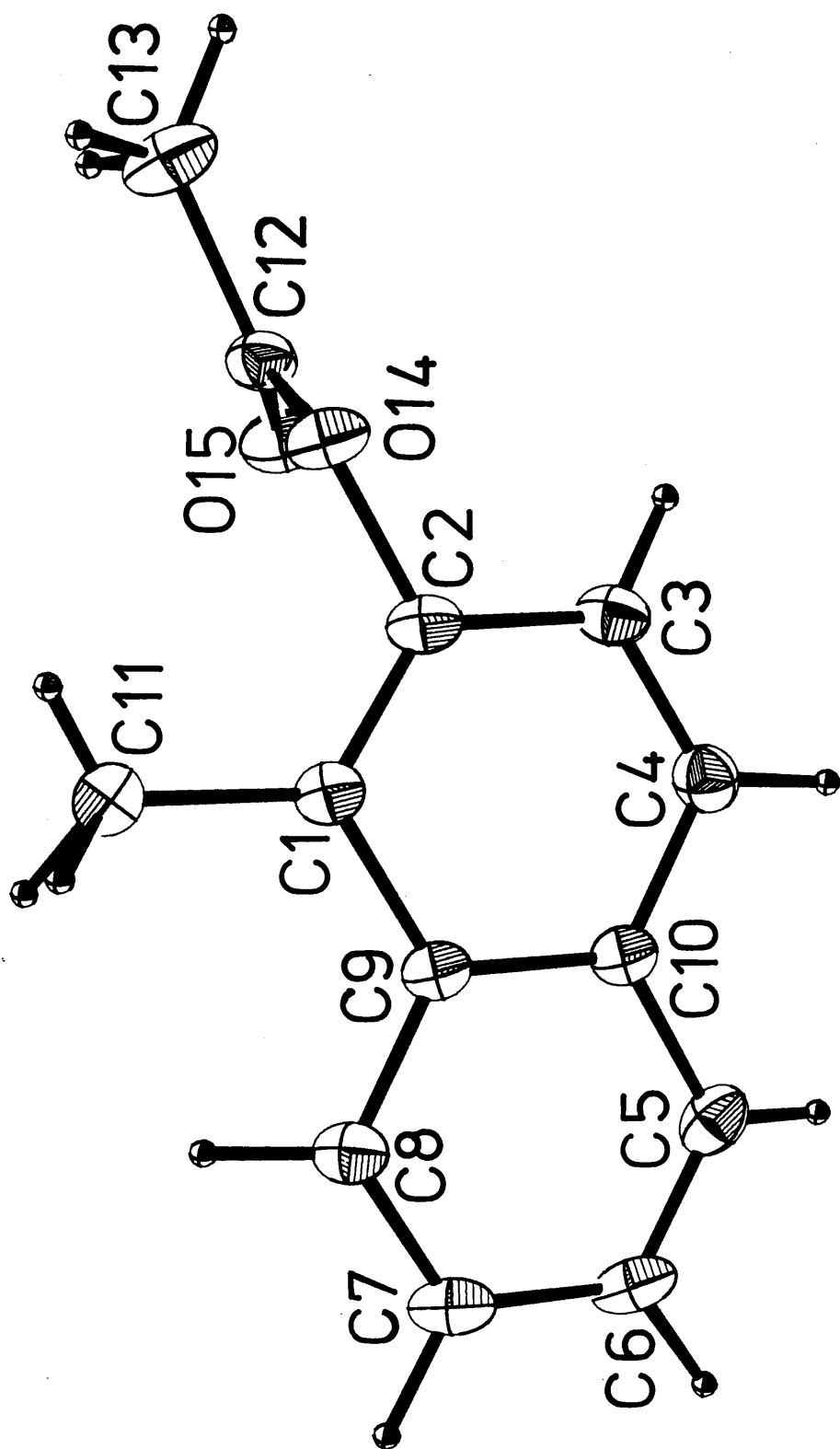
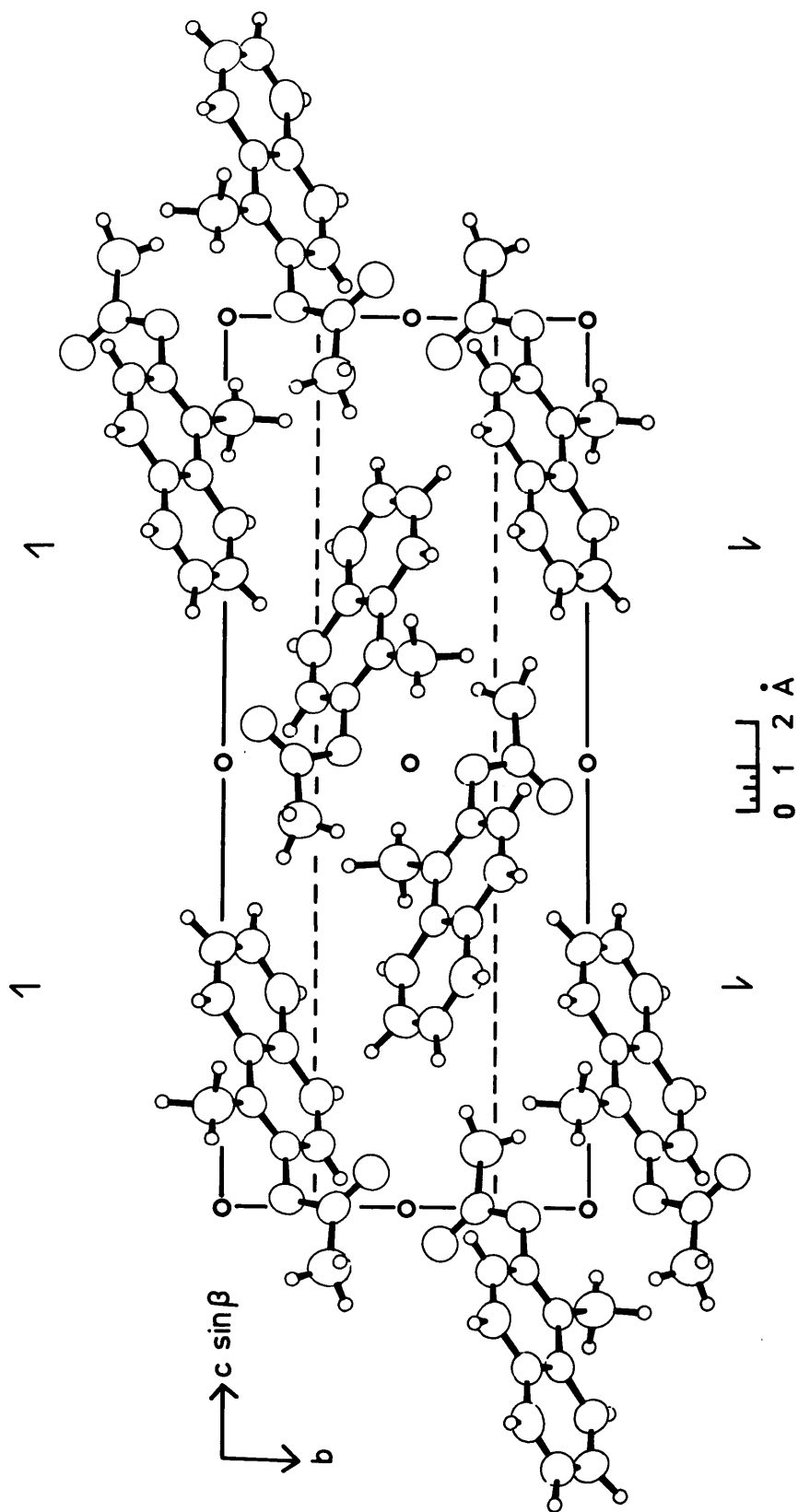


Fig. 3.3 1-Methyl-2-naphthyl acetate: Crystal packing

A view down the a axis



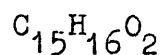


3.4 1-Isopropyl-2-naphthyl acetate

Experimental

Crystal data

1-isopropyl-2-naphthyl acetate



Unit cell dimensions \underline{a} = 19.171(5) Å°

\underline{b} = 8.591(1) Å° $\underline{\alpha} = \underline{\beta} = \underline{\gamma} = 90^\circ$

\underline{c} = 15.457(3) Å°

Space group Pbca (D_{2h}^{15})

M = 228 a.m.u.

V = 2546 Å°³

D_o = 1.07 gm.cm⁻³

D_c = 1.190 gm.cm⁻³

μ = 0.84 cm⁻¹ (Mo-K α)

$F(000)$ = 976

Z = 8

The space group was uniquely determined by the systematic absences:

hol when $l \neq 2n$, hko when $h \neq 2n$, okl when $l \neq 2n$.

The crystal used in this analysis was a diamond-shaped plate of dimensions 1x1x0.3 mm.

Data Collection

The cell dimensions were initially obtained from measurements of Weissenberg photographs. These photographs were also used to determine the space group. The crystal was then transferred to the goniometer head of a Hilger and Watts Y290 diffractometer and aligned as described previously. The cell dimensions were redetermined during this procedure and it is these values that are quoted in the crystal data.

The reflexions in the octant hkl were measured in the range $2\theta \leq 40^\circ$ using Mo-K α radiation. The following scan parameters were employed.

Increment in <u>θ</u>	0.01°
Number of scan steps	80
Time per step	1.5 sec.

Background counts of 30 seconds were taken at each end of the scan range and two strong reflexions were monitored after every 50 intensity measurements. These standard reflexions showed a significant variation towards the end of the data collection and a new set of intensity data had to be obtained using a crystal enclosed in a capillary tube. This crystal appeared to be opaque after a few days but the standards remained reasonably constant.

The measured intensities were placed on a common scale and the integrated intensities (I) yielded 2234 independant structure amplitudes a proportion of which was unobserved.

Reflexions for which $2\theta \leq 20^\circ$ were remeasured using

reduced beam intensity in order to apply a correction for counter saturation.

No correction was made for absorption.

Structure analysis

The phase problem was overcome by using the program MULTAN. The 300 reflexions with the largest values of $|E|$ ($|E| \geq 1.47$) were used to set up triplet relationships. The program selected three reflexions as origin defining and a further three for which the phase was allowed to take on values of 180° or 360° (Table X). These were then used to generate the 8 possible phase sets. The combination producing the highest combined figure of merit gave an E -map indicating all the non-hydrogen atoms.

A structure factor calculation was carried out using the 1155 independent structure amplitudes for which $|F_o| \geq 4\sigma|F_o|$ together with these atomic locations. This gave an initial R of 31.3%. On allowing these atoms to vibrate isotropically, a full matrix least squares minimisation converged at $R = 22.4\%$ after two cycles. The value of R fell to 10.0% after two further cycles of full matrix least squares during which anisotropic thermal motion was permitted.

A difference synthesis then revealed most of the hydrogen atoms. Subsequent full matrix least squares (2 cycles) with all the carbon and oxygen atoms

TABLE X SUMMARY OF STRUCTURE DETERMINATION

A. INITIAL SET OF PHASES

REFLECTION	PHASE #	MOD. E
4 1 2 +	360	3.74
2 7 3 +	360	3.23
3 6 9 +	360	2.70
3 8 3	360 OR 180	3.18
5 3 4	360 OR 180	2.95
10 2 2	360 OR 180	2.93

B. STATISTICS

	E1	E2	E3	E4	E5	E6
THEORETICAL FOR CENTRIC	0.798	1.00	0.968	31.73	4.55	0.27
THEORETICAL FOR ACENTRIC	0.886	1.00	0.736	36.79	1.83	0.01
FOUND	0.779	1.00	0.958	32.00	4.60	0.4

* IN DEGREES
+ ORIGIN DEFINING

MOD. E	MODULUS OF E FOR REFLECTION CONCERNED
E1	MEAN VALUE FOR MODULUS OF E
E2	MEAN VALUE FOR MODULUS OF E**2
E3	MEAN VALUE FOR MODULUS OF (E**2)-1)
E4	% OF REFLECTIONS WITH MODULUS OF E > 1.0
E5	% OF REFLECTIONS WITH MODULUS OF E > 2.0
E6	% OF REFLECTIONS WITH MODULUS OF E > 3.0

(anisotropic motion), these hydrogen positions (isotropic vibration), and also including an allowance for counter saturation reduced \underline{R} to 8.5% .

The remaining hydrogen coordinates were obtained from a new difference Fourier summation. A weighting scheme of the form:

$$W = 1/(A + BF_o + CF_o^2) \quad \text{with} \quad \begin{array}{l} A = 0.4999 \\ B = -0.3020 \\ C = 0.0049 \end{array}$$

was introduced and two further cycles of full matrix least squares calculations produced a final \underline{R} of 6.1% .

This structure determination was carried out in collaboration with Dr. P. J. Mallinson (University of Glasgow) who collected the data and solved the structure; the author being responsible for the subsequent refinement.

3.5 Results

Table XI 1-Isopropyl-2-naphthyl acetate: Fractional atomic
coordinates

Positional estimated standard deviations are
shown in parentheses.

The hydrogen atoms are numbered according to the
atoms to which they are attached.

The table shows:

Atom	x	y	z
------	---	---	---

C(1)	1432(3)	8663(5)	2290(3)
C(2)	0908(3)	8983(6)	1715(3)
C(3)	0469(3)	10296(6)	1763(3)
C(4)	0562(3)	11325(6)	2412(4)
C(5)	1163(3)	12127(6)	3755(4)
C(6)	1646(3)	11869(7)	4372(3)
C(7)	2088(3)	10589(7)	4314(4)
C(8)	2029(3)	9574(6)	3648(3)
C(9)	1518(3)	9745(6)	2988(3)
C(10)	1083(3)	11088(6)	3049(3)
C(11)	1879(2)	7193(6)	2222(3)
C(12)	0740(3)	8276(7)	0231(3)
C(13)	0470(3)	6958(7)	-0301(3)
C(17)	1455(4)	5729(7)	2396(5)
C(18)	2274(3)	7090(7)	1374(4)
O(14)	0744(2)	7858(4)	1083(2)
O(15)	0929(2)	9496(5)	-0026(3)
H(3)	0114(26)	10370(61)	1253(32)
H(4)	0266(24)	12342(61)	2424(31)
H(5)	0855(23)	13030(53)	3719(30)
H(6)	1686(24)	12616(61)	4874(29)
H(7)	2421(24)	10459(51)	4729(26)
H(8)	2325(20)	8747(49)	3601(25)
H(11)	2265(27)	7276(65)	2686(33)
H(131)	0534(37)	7043(92)	-0963(48)
H(132)	-0024(31)	6946(66)	-0308(36)
H(133)	0670(41)	6045(79)	-0150(48)
H(171)	1750(26)	4924(58)	2401(33)
H(172)	1229(38)	5782(88)	2949(54)
H(173)	1067(33)	5577(76)	1932(42)
H(181)	2663(26)	6203(64)	1419(31)
H(182)	2494(28)	7935(62)	1210(34)
H(183)	1936(42)	6796(84)	0677(44)

Table XII 1-Isopropyl-2-naphthyl acetate: Thermal parameters

Carbon and oxygen values are anisotropic.

Hydrogen values are isotropic.

Estimated errors in the last digit are shown
in parentheses.

All figures are multiplied by 10^3 .

The anisotropic temperature factor expression
used was of the form described by equation (12)
in chapter one, the coefficients being in terms
of the mean square amplitudes of vibration U_{ij} .

The table shows:

Atom	U_{11} or U	U_{22}	U_{33}	U_{13}	U_{12}	U_{23}
------	-----------------	----------	----------	----------	----------	----------

C(1)	49(3)	43(3)	44(3)	4(3)	8(3)	8(2)
C(2)	49(3)	40(3)	46(3)	-5(3)	3(3)	3(3)
C(3)	51(3)	55(3)	53(3)	14(3)	-1(3)	3(3)
C(4)	57(3)	45(3)	64(4)	19(3)	3(3)	6(3)
C(5)	64(4)	46(3)	64(4)	-1(3)	18(3)	2(3)
C(6)	72(4)	59(4)	53(3)	-12(3)	2(3)	-6(3)
C(7)	63(4)	64(4)	57(4)	-8(3)	1(3)	1(3)
C(8)	47(3)	48(3)	68(3)	5(3)	1(3)	5(3)
C(9)	43(3)	51(3)	44(3)	-1(3)	8(2)	11(2)
C(10)	49(3)	39(3)	54(3)	2(3)	8(3)	8(3)
C(11)	50(3)	45(3)	54(3)	9(3)	7(3)	8(3)
C(12)	57(3)	56(4)	53(4)	5(3)	-3(3)	5(3)
C(13)	75(4)	67(4)	58(3)	11(3)	-9(3)	-12(3)
C(17)	78(4)	51(4)	118(6)	14(3)	17(4)	30(4)
C(18)	71(4)	57(4)	68(4)	19(3)	21(3)	8(3)
O(14)	59(2)	43(2)	51(2)	-3(2)	-4(2)	6(2)
O(15)	127(4)	66(3)	59(2)	-22(3)	3(3)	16(2)
H(3)	70(16)					
H(4)	62(15)					
H(5)	52(14)					
H(6)	59(15)					
H(7)	47(14)					
H(8)	33(12)					
H(11)	78(18)					
H(131)	136(29)					
H(132)	87(19)					
H(133)	130(28)					
H(171)	63(16)					
H(172)	152(30)					
H(173)	108(22)					
H(181)	63(16)					
H(182)	75(17)					
H(183)	138(28)					

TABLE XIII

1-ISOPROPYL-2-NAPHTHYL-ACETATE:
INTRAMOLECULAR BONDED DISTANCES (ANGSTROMS)
STANDARD DEVIATIONS (ANGSTROMS) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	DIST.	ATOM1	ATOM2	DIST.
C1	C2	1.371(7)	C12	O15	1.179(7)
C1	C9	1.431(7)	C3	H3	1.041(53)
C1	C11	1.524(7)	C4	H4	1.033(53)
C2	C3	1.408(7)	C5	H5	0.976(47)
C2	O14	1.414(6)	C6	H6	1.017(51)
C3	C4	1.348(8)	C7	H7	0.910(46)
C4	C10	1.412(8)	C8	H8	0.920(43)
C5	C6	1.351(8)	C11	H11	1.052(55)
C5	C10	1.420(8)	C13	H131	1.046(77)
C6	C7	1.390(9)	C13	H132	0.933(64)
C7	C8	1.353(8)	C13	H133	0.942(75)
C8	C9	1.428(7)	C17	H171	0.892(53)
C9	C10	1.430(7)	C17	H172	0.977(88)
C11	C17	1.522(8)	C17	H173	1.038(69)
C11	C18	1.518(8)	C18	H181	1.023(55)
C12	C13	1.492(8)	C18	H182	0.891(56)
C12	O14	1.367(6)	C18	H183	1.042(80)

TABLE XIV 1-ISOPROPYL-2-NAPHTHYL-ACETATE: SOME INTRAMOLECULAR
NON-BONDED SEPARATIONS

ATOM1	ATOM2	DIST #	ATOM1	ATOM2	DIST #
C(1).....C(8)		2.516	O(15).....H(3)		2.63
C(11).....O(14)		2.855	O(15).....H(183)		3.32
C(11).....H(8)		2.657	H(4).....H(5)		2.37
O(14).....C(17)		3.053	H(7).....H(8)		2.29
O(14).....C(18)		3.039	H(8).....H(11)		1.90

IN ANGSTROMS

TABLE XV 1-ISOPROPYL-2-NAPHTHYL ACETATE: INTERMOLECULAR CONTACTS OF LESS THAN 4.0 ANGSTROMS

ATOM1	ATOM2	DIST #	ATOM1	ATOM2	DIST #
C(5).....C(17)	*	3.782	C(4).....C(17)	***	3.910
C(13).....C(13)	**	3.917	C(4).....O(14)	***	3.662
C(6).....C(8)	***	3.619	C(5).....C(13)	***	3.940
C(7).....C(8)	***	3.955	C(5).....O(14)	***	3.718
C(8).....C(11)	***	3.783	C(7).....C(13)	***	3.839
C(8).....C(17)	***	3.630	C(7).....C(18)	***	3.945
C(9).....C(11)	***	3.908	C(8).....C(13)	***	3.645
C(9).....C(18)	***	3.956	C(9).....C(13)	***	3.627
C(3).....C(17)	***	3.927	C(10).....C(13)	***	3.834
C(4).....C(2)	***	3.869			

IN ANGSTROMS

THE ASTERISKS(*) REFER TO THE FOLLOWING TRANSFORMATIONS OF THE ATOMIC COORDINATES

*	X, 1.0+Y, Z
**	-X, 1.0-Y, -Z
***	0.5-X, 0.5+Y, Z
****	-X, 0.5+Y, 0.5-Z
*****	X, 1.5-Y, 0.5+Z

TABLE XVI

1-ISOPROPYL-2-NAPHTHYL-ACETATE: VALENCY ANGLES(DEGREES)
STANDARD DEVIATIONS(DEGREES) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	ATOM3	ANGLE	ATOM1	ATOM2	ATOM3	ANGLE
C9	C1	C2	115.7(4)	C11	C1	C2	122.2(4)
C11	C1	C9	122.1(4)	O14	C2	C1	118.2(4)
C3	C2	C1	124.7(5)	O14	C2	C3	116.8(4)
C4	C3	C2	119.0(5)	C10	C4	C3	121.1(5)
C10	C5	C6	121.0(5)	C7	C6	C5	120.2(5)
C8	C7	C6	120.7(5)	C9	C8	C7	122.4(5)
C10	C9	C1	120.9(4)	C8	C9	C1	123.2(5)
C10	C9	C8	115.9(4)	C5	C10	C4	121.6(5)
C9	C10	C4	118.6(4)	C9	C10	C5	119.8(5)
C18	C11	C1	112.9(4)	C17	C11	C1	111.7(4)
C18	C11	C17	112.2(5)	O14	C12	C13	109.5(4)
O15	C12	C13	127.0(5)	O15	C12	O14	123.4(5)
C12	O14	C2	119.7(4)	H3	C3	C2	113.8(31)
H3	C3	C4	127.2(31)	H4	C4	C3	119.3(28)
H4	C4	C10	119.4(28)	H5	C5	C6	126.0(28)
H5	C5	C10	112.9(28)	H6	C6	C5	119.7(29)
H6	C6	C7	120.1(29)	H7	C7	C6	118.0(29)
H7	C7	C8	121.3(29)	H8	C8	C7	121.8(26)
H8	C8	C9	115.8(26)	H11	C11	C1	107.1(32)
H11	C11	C17	108.4(31)	H11	C11	C18	104.0(30)
H131	C13	C12	115.9(44)	H132	C13	C12	110.0(37)
H133	C13	C12	111.9(47)	H132	C13	H131	96.5(54)
H133	C13	H131	104.3(64)	H133	C13	H132	117.6(62)
H172	C17	C11	111.8(47)	H173	C17	C11	111.0(38)
H171	C17	C11	106.7(34)	H172	C17	H171	107.7(59)

VALENCY_ANGLES (CONTINUED)

ATOM1	ATOM2	ATOM3	ANGLE	ATOM1	ATOM2	ATOM3	ANGLE
H173	C17	H171	110.0(51)	H173	C17	H172	109.6(60)
H181	C18	C11	109.7(29)	H182	C18	C11	115.1(36)
H183	C18	C11	107.8(45)	H182	C18	H181	104.8(47)
H183	C18	H181	110.5(54)	H183	C18	H182	109.0(56)

TABLE XVII

1-10PROPXYL-2-NAPHTHYL-ACETATE: TORSION ANGLES(DEGREES)
STANDARD DEVIATIONS(DEGREES) ARE SHOWN IN PARENTHESES

ATOM1	ATOM2	ATOM3	ATOM4	ANGLE	ATOM1	ATOM2	ATOM3	ATOM4	ANGLE
C9	C1	C2	C3	1.7(8)	C9	C1	C2	O14	-172.1(4)
C11	C1	C2	C3	178.3(5)	C11	C1	C2	O14	4.5(7)
C2	C1	C9	C8	178.8(5)	C2	C1	C9	C10	-1.8(7)
C11	C1	C9	C8	2.1(8)	C11	C1	C9	C10	-178.4(4)
C2	C1	C11	C17	-67.0(7)	C2	C1	C11	C18	60.5(7)
C9	C1	C11	C17	109.4(6)	C9	C1	C11	C18	-123.1(5)
C1	C2	C3	C4	-0.4(8)	O14	C2	C3	C4	173.5(5)
C1	C2	O14	C12	-125.9(5)	C3	C2	O14	C12	59.9(6)
C2	C3	C4	C10	-0.9(8)	C3	C4	C10	C5	-177.0(5)
C3	C4	C10	C9	0.7(8)	C10	C5	C6	C7	1.5(9)
C6	C5	C10	C4	178.0(5)	C6	C5	C10	C9	0.3(8)
C5	C6	C7	C8	-1.5(9)	C6	C7	C8	C9	-0.4(9)
C7	C8	C9	C1	-178.4(5)	C7	C8	C9	C10	2.1(8)
C1	C9	C10	C4	0.7(8)	C1	C9	C10	C5	178.5(5)
C8	C9	C10	C4	-179.9(5)	C8	C9	C10	C5	-2.0(7)
C13	C12	O14	C2	-173.4(4)	O15	C12	O14	C2	7.6(8)
C2	C1	C11	H11	174.3(32)	C9	C1	C11	H11	-9.2(32)
C1	C2	C3	H3	177.3(33)	O14	C2	C3	H3	-8.9(33)
C2	C3	C4	H4	174.6(33)	H3	C3	C4	C10	-178.2(38)
H3	C3	C4	H4	-2.6(50)	H4	C4	C10	C5	7.4(34)
H4	C4	C10	C9	-174.8(33)	C10	C5	C6	H6	-179.5(33)
H6	C5	C6	C7	-174.8(35)	H5	C5	C6	H6	4.2(48)
H5	C5	C10	C4	-5.2(31)	H5	C5	C10	C9	177.0(30)
C6	C6	C7	H7	177.6(33)	H6	C6	C7	C8	179.5(33)
H6	C6	C7	H7	-1.3(47)	C6	C7	C8	H8	177.6(31)

DIEDION_ANGLES (CONTINUED)

ATOM1	ATOM2	ATOM3	ATOM4	ANGLE	ATOM1	ATOM2	ATOM3	ATOM4	ANGLE
H7	C7	C8	C9	-179.5(34)	H7	C7	C8	H8	-1.5(47)
H8	C8	C9	C1	3.5(30)	H8	C8	C9	C10	-176.0(29)
C1	C11	C17	H171	-174.1(36)	C1	C11	C17	H172	-56.6(52)
C1	C11	C17	H173	66.1(41)	C18	C11	C17	H171	58.0(36)
C18	C11	C17	H172	175.6(52)	C18	C11	C17	H173	-61.8(41)
H11	C11	C17	H171	-56.2(48)	H11	C11	C17	H172	61.3(62)
H11	C11	C17	H173	-176.0(52)	C1	C11	C18	H181	164.6(32)
C1	C11	C18	H182	46.8(40)	C1	C11	C18	H183	-75.1(46)
C17	C11	C18	H181	-68.1(32)	C17	C11	C18	H182	174.1(40)
C17	C11	C18	H183	52.2(46)	H11	C11	C18	H181	48.8(45)
H11	C11	C18	H182	-69.0(51)	H11	C11	C18	H183	169.2(56)
O14	C12	C13	H131	-167.3(47)	O14	C12	C13	H132	84.7(40)
O14	C12	C13	H133	-47.9(51)	O15	C12	C13	H131	11.6(47)
O15	C12	C13	H132	-96.4(41)	O15	C12	C13	H133	131.0(51)

THE ANGLE 1-2-3-4 IS DEFINED AS POSITIVE IF, WHEN VIEWED ALONG THE 2-3 BOND, ATOM 1 HAS TO BE ROTATED CLOCKWISE TO ECLIPSE ATOM 4

TABLE XVIII 1-ISOPROPYL-2-NAPHTHYL-ACETATE: MEAN PLANE CALCULATIONS

ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
1.		
C(1)		-0.002(5)
C(2)		0.024(5)
C(3)		0.017(5)
C(4)		-0.019(5)
C(5)		-0.002(5)
C(6)		0.030(6)
C(7)		0.010(6)
C(8)		-0.017(5)
C(9)		-0.013(5)
C(10)		-0.026(5)
	C(11)	0.049(5)
	O(14)	0.205(3)
	H(4)	-0.102(49)
	H(5)	-0.058(45)
	H(8)	-0.051(39)

TABLE XVIII (CONTINUED)

	ATOMS IN PLANE	ATOMS OUT OF PLANE	DEVIATION (ANGSTROMS)
2.	C (12) C (13) O (14)		0.000 (5) 0.000 (6) 0.000 (3)
		O (15)	-0.013 (5)

THE OUT OF PLANE ATOMS WERE NOT INCLUDED IN THE DERIVATION OF THE PLANES.

THE EQUATIONS OF THESE PLANES ARE:

1. $-0.6516X - 0.5278Y + 0.5448Z = -3.7867$
2. $-0.9315X + 0.3493Y + 0.1015Z = 1.1986$

Fig. 3.4 1-Isopropyl-2-naphthyl acetate: A view of the molecule



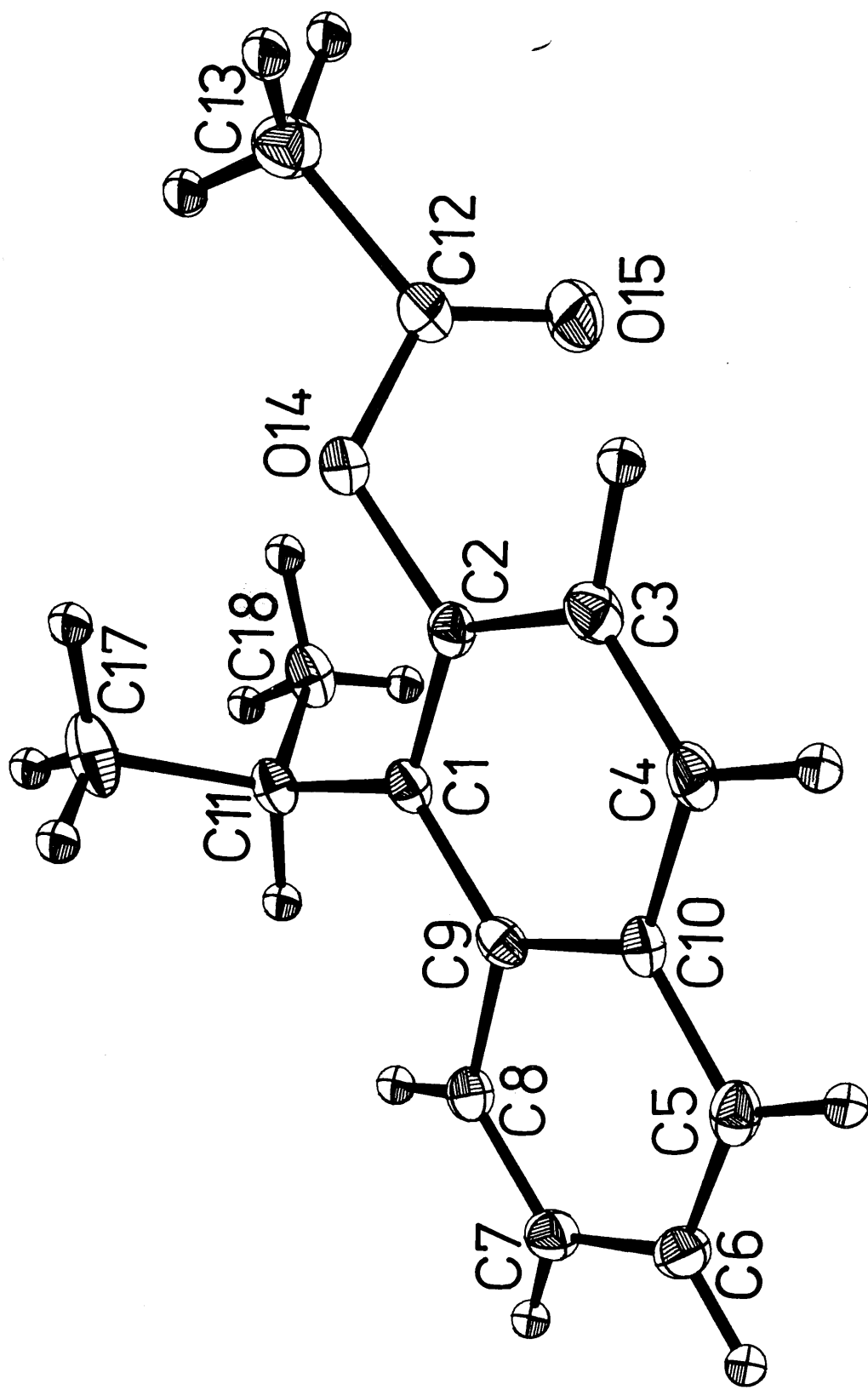
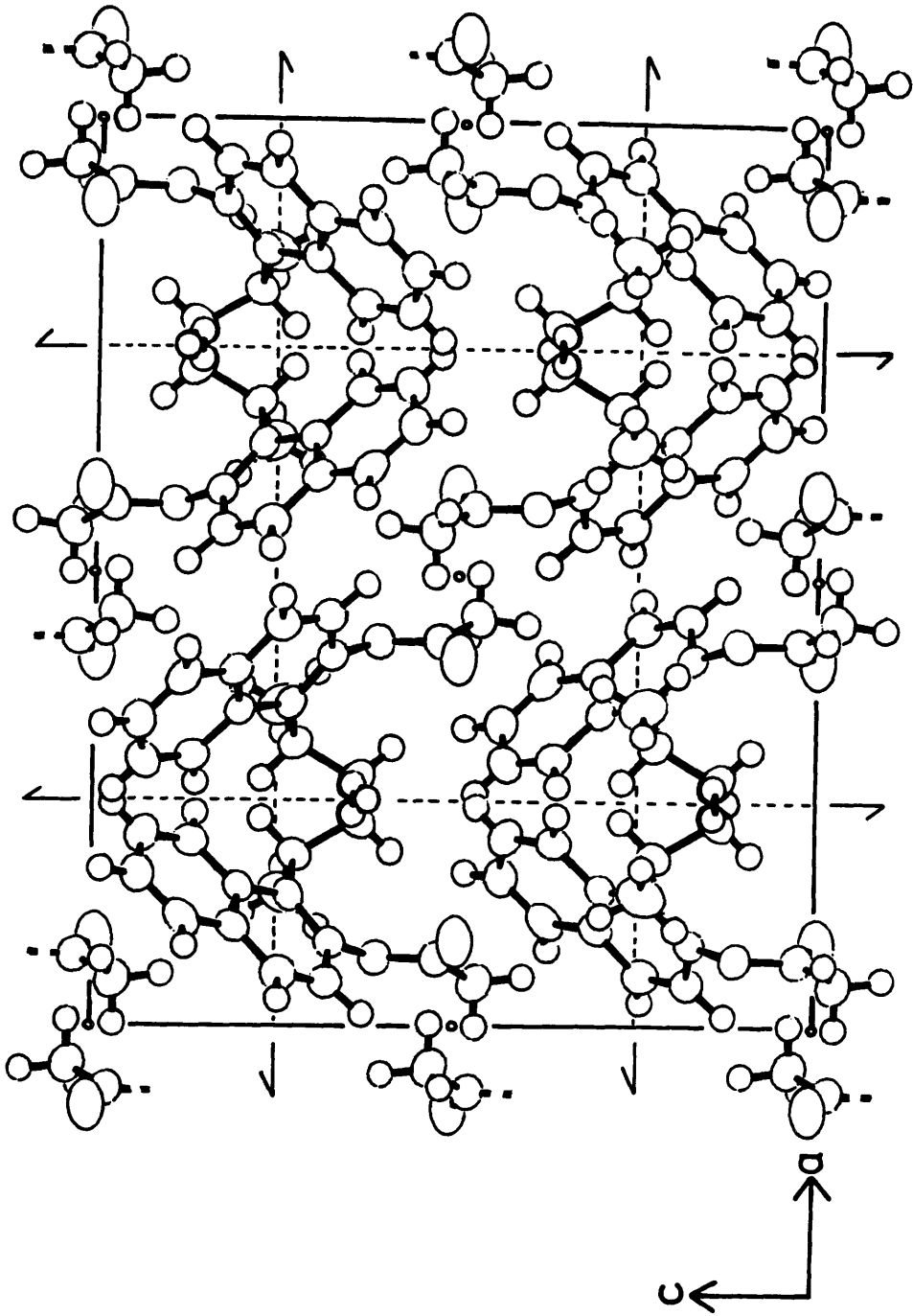


Fig. 3.5 1-Isopropyl-2-naphthyl acetate: Crystal packing

A view down the b axis



0 1 2 Å

3.6 Discussion of results

Tables II - IX and XI - XVIII summarise the results of the analyses for the methyl- and isopropyl- derivatives of 2-naphthyl acetate respectively. Figures 3.2 and 3.4 illustrate views of the respective molecules and Fig. 3.3 shows the crystal packing of 1-methyl-2-naphthyl acetate viewed along the a axis while Fig. 3.5 is a drawing of the crystal packing of its isopropyl homologue viewed along the b axis.

The final and observed structure factors are listed in Tables XX and XXI respectively.

The analysis shows that the geometry of the methyl compound is very similar to that observed in naphthalene (3)

Fig. 3.6 Structure of methyl compound

Internal values	Bond lengths (Å)
Apical values	Valence angles (°)
External values	Torsion angles (°)

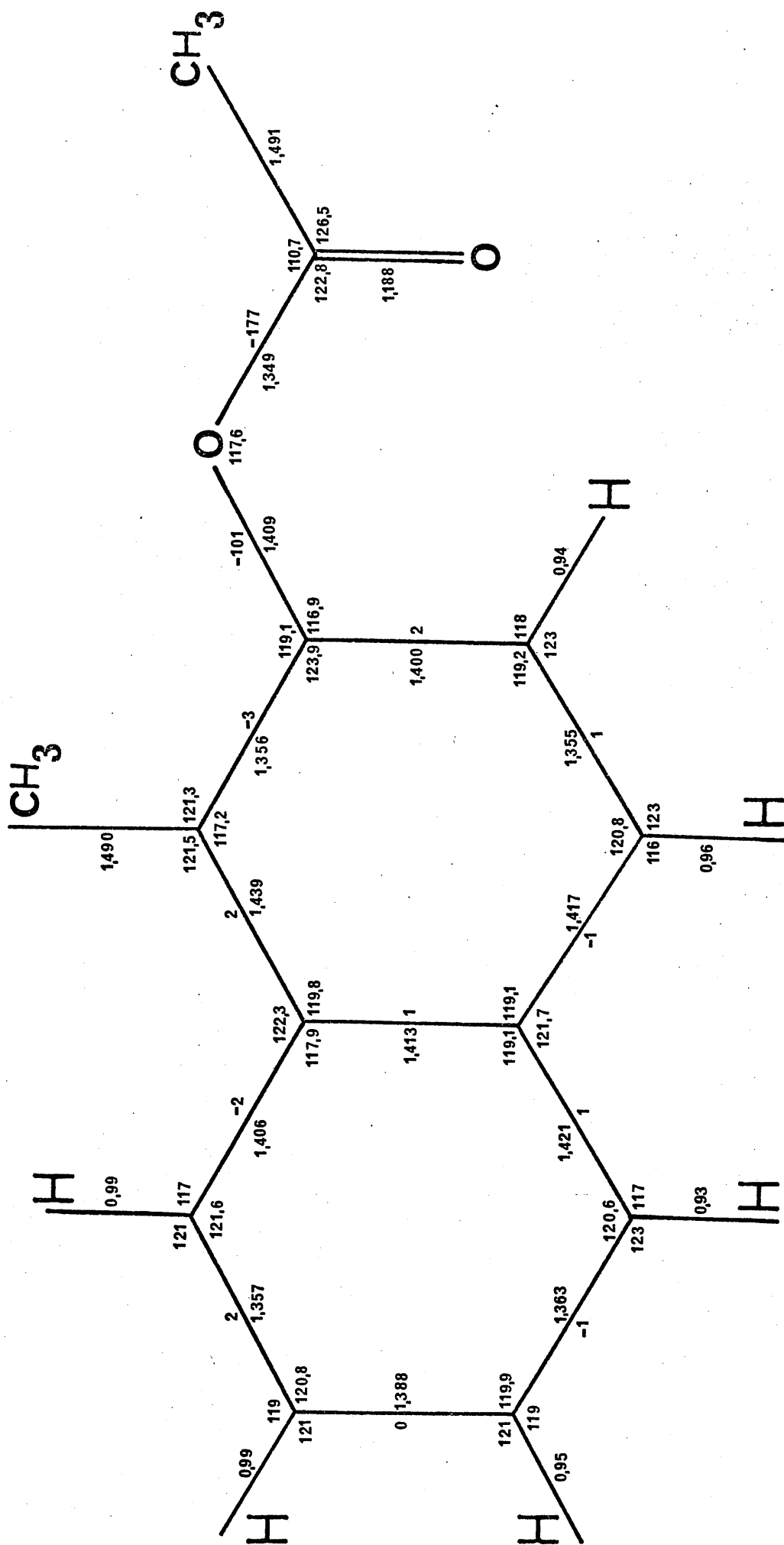




Fig. 3.7 Structure of isopropyl compound

Internal values

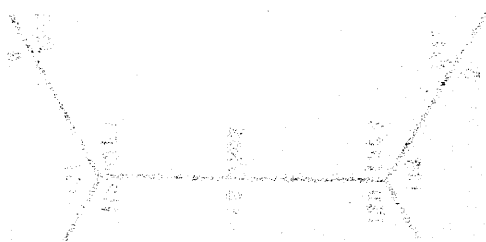
Bond lengths (\AA)

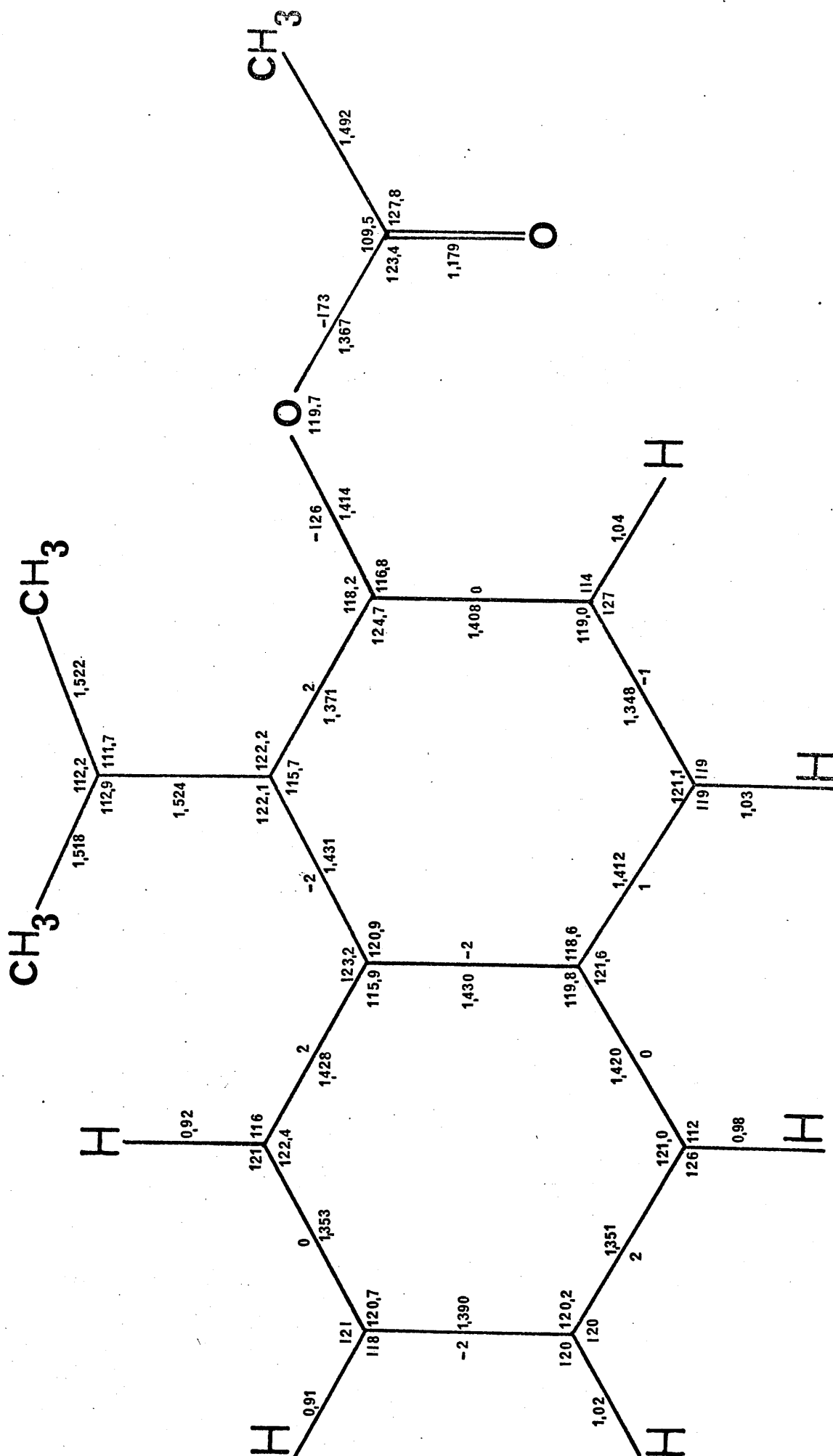
Apical values

Valence angles ($^\circ$)

External values

Torsion angles ($^\circ$)





indicating that very little additional strain has been imposed by the 1, 2- substituents. The naphthalene nucleus is essentially planar since the maximum ring torsion angle (ω) is -3° with $|\omega| = 16^\circ$ and $\omega = 2$. The largest displacement from the mean plane of the ring is 0.03\AA at C(2). (Table IX). This is not entirely unexpected because previous work has shown that peri-strain is usually minimized by out of plane bending and in-plane splaying of the 1, 8- substituents and that the naphthalene system is only distorted in extreme cases. Thus, 1, 8- dimethyl-naphthalene (4) and 1, 8- diphenyl-naphthalene (5) have planar aromatic systems, while 1, 8- di(bromomethyl)-naphthalene (6); 1, 4, 5, 8, -tetrachloronaphthalene (7) and 1, 4, 5, 8- tetraphenyl-naphthalene (8) all exhibit deformation of the naphthalene nucleus.

Any peri -strain present in 1- methyl-2-naphthyl acetate must therefore manifest itself by causing either out of plane or in-plane deformations. The out of plane bending of C(11) and H(8) is minimal as the displacements of these atoms from the mean plane of the naphthalene ring (-0.071 and 0.02\AA respectively) are barely significant although they may be in the opposite sense. Likewise, the C(11) - C(1) - C(9) - C(8) torsion angle of 3° , the C(1) - C(9) - C(8) - H(8) value of 2° and the angle between the planes defined by C(1) - C(8) - H(8) and C(8) - C(1) - C(11), (4°), hardly provide reasonable evidence of out of plane bending.

The evidence for in-plane splaying is more satisfactory. The C(1) - C(9) - C(8) valence angle opens up to 122.3° , which is over twenty standard deviations

greater than the "equilibrium" value of 120° , and 1.0° larger than that observed in unsubstituted naphthalene (3). The C(9) - C(1) - C(11) angle is 121.5° and this also indicates some in-plane deformation. There is a fairly large estimated standard deviation (1.5°) on the H(8) - C(8) - C(9) value of 117° so that it is difficult to assess the distortion at C(8) from this measurement. However, the C(1) C(8) intramolecular non-bonded distance of 2.494\AA is a little longer than the C(4).... C(5) separation (2.481\AA) and this difference is just large enough to indicate the necessity for some relief of strain at C(1), C(8), but comparison with the geometry in this region in 1, 8 disubstituted naphthalenes (e.g. 4, 6) shows the relatively small amount of strain present in 1-methyl - 2 - naphthylacetate.

There are also variations in the naphthalene nucleus H-C-C bond angles but these angles have standard deviations which are too large for them to be used to investigate the possible transmission of peri-strain around the ring as reported by Bright et al (4) but the H(4).....H(5) separation of 2.39\AA is reasonably correct and not shortened to c. 2.2\AA as in 1, 8 dimethyl naphthalene so that this transmission effect is not observed.

The previous discussion shows that 1-methyl-2-naphthyl acetate is far less strained than the 1, 8 disubstituted naphthalenes for which results are available. The rate measurements on the autoxidation of alkyl naphthols by Brady and Carnduff (2) suggest that the

replacement of the methyl substituent at C(1) by an isopropyl function should cause a substantial increase in peri-strain.

The present study does not support this suggestion. A comparison of the pertinent geometrical parameters for the two compounds is given in Table XIX. In general the increase in strain in going from C(1)-methyl to C(1)-isopropyl is not as large as might be anticipated from the rate measurements.

Thus the relative values for the torsion angles and out of plane displacements in the naphthalene nucleus are unchanged within the limits of this experiment as are the displacements of C(11) and C(8) from the mean plane of the aromatic system.

Out of plane deformation of C(11) and H(8) is not increased in the isopropyl homologue since neither the displacement of these atoms from the mean plane of the naphthalene ring nor the C(11) - C(1) - C(9) - C(8) and C(1) - C(9) - C(8) - H(8) torsion angles show any significant variation. The angle between the planes defined by C(11), C(2), C(8) and H(8), C(8), C(1) remains at 4° .

In-plane bending at C(1) and C(8) is again the only type of distortion which is large enough to be significant.

The isopropyl substitution for methyl at C(1) opens the C(11) - C(1) - C(9) angle by a further 0.6° to 122.1° and the C(1) - C(9) - C(8) value increases by almost 1°

TABLE XIX COMPARISON OF PERI-SUBSTITUENT IN METHYL AND ISOPROPYL DERIVATIVES

PARAMETER	METHYL	ISOPROPYL
MAX. TORSION ANGLE IN NAPHTHALENE NUCLEUS *	3	2
OMEGA	2	0
MOD. OMEGA	16	14
MAX. DISPLACEMENT OF RING CARBON ATOMS ++	0.03 AT C(2)	0.03 AT C(6)
MAX. DISPLACEMENT OF C(11) ++	-0.07	0.05
H(8) ++	0.02	-0.05
NON-BONDED DISTANCES +		
C(1)...C(8)	2.494	2.516
C(4)...C(5)	2.481	2.472
O(15)...H(3)	3.18	2.63
H(4)...H(5)	2.39	2.37
VALENCE ANGLES *		
C(9)-C(1)-C(11)	121.5	122.1
C(1)-C(9)-C(8)	122.3	123.2
C(4)-C(10)-C(5)	121.8	121.6
C(9)-C(8)-H(8)	117	116
TORSION ANGLES *		
C(11)-C(1)-C(9)-C(8)	3	2
C(1)-C(2)-O(14)-C(12)	-101	-126
C(1)-C(9)-C(8)-H(8)	2	4
"C(11)-C(1)-C(8)-H(8)"	4	4

SYMBOLS EXPLAINED ON FOLLOWING PAGE

EXPLANATION OF SYMBOLS IN TABLE XIX

IN DEGREES
IN ANGSTROMS

OMEGA

MOD. OMEGA

THE SUM OF THE TORSION ANGLES OF THE NAPHTHALENE NUCLEUS

THE SUM OF THE MODULII OF THE TORSION ANGLES OF THE NAPHTHALENE NUCLEUS

FROM MEAN PLANE THROUGH NAPHTHALENE NUCLEUS (ANGSTROMS)

* +

+

to 123.2° . Similar enlargement of the latter angle has also been observed in 1,8-dimethyl-naphthalene (4) (125.2°) and in 1,8-di(bromomethyl)-naphthalene (6) (127°) so there is clearly a progression in the distortion of C(1) - C(9) - C(8) from 122.3° (1-methyl) to 127° (1,8-di(bromomethyl)).

This deformation increases the C(1) C(8) separation throughout the series. The relevant values being 2.494\AA° (methyl), 2.516\AA° (isopropyl), 2.543 (1,8-dimethyl), and 2.57 (1,8-di(bromomethyl)).

1-Isopropyl-2-naphthyl acetate shows no evidence for transmission of the peri-strain to the C(4), C(5) locations since neither the C(4)C(5) nor H(4) H(5) distances (2.472\AA° and 2.37\AA° respectively) are decreased by substantial amounts.

In the isopropyl derivative the C(11) substituents are arranged so that H(11) almost eclipses the C(1) - C(9) bond (C(9) - C(1) - C(11) - H(11) torsion angle is -9°) this is in contrast to the situation in the methyl compound where one of the C(11) hydrogens is located so as to give a C(2) - C(1) - C(11) - H(112) torsion angle of 2° . H(11) appears to be separated from H(8) by 1.90\AA° (Table XIV) but the estimated standard deviations of the fractional co-ordinates of these atoms are in the range 0.002 to 0.006\AA° and the C(8) - H(8) bond length is found as 0.92\AA° (Table XIII) so that this non-banded distance is probably not significant.

Further evidence of a difference in strain in these two 1-alkyl-2-naphthyl acetates is provided by the conformation adopted by the acetate side chain. In the methyl derivative the ester function rotates with respect to the naphthalene nucleus to give a C(1) - C(2) - O(14) - C(12) torsion angle of -101° . The reason for this is to reduce the H(3) O(15) non-bonded interaction by increasing the separation of these two atoms to 3.18\AA . In the isopropyl compound the replacement of two of the C(11) hydrogens by methyl functions restricts the amount of torsional rotation in the acetate group and the C(1) - C(2) - O(14) - C(12) torsion angle is then -126° . This is only sufficient to separate H(3) and O(15) by 2.63\AA .

The results from the x-ray analysis of these two derivatives of 2-naphthol do not confirm the hypothesis of Brady and Carnduff regarding steric strain in 1-alkyl-2-naphthols. Subsequent crystal analysis on other compounds with different substituents at R1, R2 and R3 (Fig. 3.1) when compared with this study may allow an interpretation of the observed rates of autoxidation.

Table XX 1-Methyl-2-naphthyl acetate: Final structure factors

The table shows:

L^*	$ F_o $	$ F_c $	Phase($^\circ$)
-------	---------	---------	-------------------

* Reflections sorted into groups with common
H,K indices (shown in each group heading)

#

7,4,L

0	13	8	-5	12	9	7	16	13	-5	17	18	12	13	7	-6	134	-130	4	23	21
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6,3,L

6,0,L

5,6,L

7,3,L

6,5,L

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6,1,L

7,2,L

6,4,L

5,8,L

5,5,L

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-3 26 26

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-17 39 -43
-18 46 -53
-19 51 53
22 17 15
20 77 75
18 34 31
16 28 26
14 16 19
12 45 42
10 20 16
6 102 106
4 75 73
2 134 135
0 251-258
-2 218 217
-4 102 102
-6 169 166
-8 30 -31
-10 126 125
-12 33 -27
-14 42 -33
-16 185-196
-18 27 -32
-20 14 -18

Table XXI 1-Isopropyl-2-naphthyl acetate: Final structure factors

The table shows:

H^*	$ F_o $	$ F_c $	Phase($^\circ$)
-------	---------	---------	-------------------

* Reflections sorted into groups with common
K,L indices (shown in each group heading)

Values of $|F_o|$ with the suffix L are
"unobserved".

[illegible]

[illegible]

[illegible]

1	192	-192	10	50L	-64	133	1	195	178	14	126	-129	5	47L	16	192	-192
2	293	-291	11	123	-64	133	2	139	-134	15	41L	-129	6	56	57	291	-82
3	54	44	12	257	-257	-208	3	17L	-20	16	0L	-5	7	275	24	177	177
4	178	153	13	196	-208	-40	4	100	-95	17	0L	3	8	22L	67	182	182
5	725	715	14	22L	-40	-11	5	136	-141	18	77	-66	9	182	61	164	164
6	246	-249	15	44L	-11	-12	6	0L	44	19	0L	-44	10	67	53L	-46	-46
7	134	-162	16	47L	40	-48	7	102	119	12	18L	1	11	164	52L	48	48
8	58	64	17	32L	-48	0	8	181	-190	13	77	-66	12	53L	67	-13	-13
9	145	-144	18	41L	0	12	9	104	106	14	0L	-44	13	53L	17L	-25	-25
10	33L	-35	19	0L	12		10	19L	-13	15	238	-207	14	53L	31L	-41	-41
11	208	-219	20				11	249	-261	16	44L	24	15	17L	49L	12	12
12	279	-287					12	159	162	17	238	249	16	49L			
13	115	-110					13	87	92	18	0L	12	17				
14	248	253					14	61	52	19	35L	73	18				
15	100	91					15	0L	-12	20	65	-74	19				
16	114	108					16	49L	-7	21	191	205	20				
17	66	28					17	0L	-41	22	256	264	21				
18	12L	4					18	39L	7	23	82	-82	22				
19	23L	39					19	38L	31	24	162	160	23				

[illegible]

15 20L -13

17 0L 47

1999-2000

三

101-7

084-1

192-454

497-54

317 33

10 166 181

12 21L -59

14 72 82

16 57L 48

-52-

70

494

1	143	154
2	153	164

4000

[illegible]

[illegible]

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PART II

MOLECULAR MECHANICS

CHAPTER FOUR

Molecular mechanics: An introduction

4.1 Introduction

Much of the progress in organic chemistry in the last 100 years has in some way been connected with an increased understanding of the spatial requirements of molecules. It has also become clear that the chemistry of a compound is very much related to its structure and hence its "structural energy". A knowledge of these is quite clearly very desirable when forecasting the outcome of chemical reactions and particularly when allied to similar information regarding possible intermediates. These various differences in reaction rates, pathways and products have been explained qualitatively by the principle of 'tendency towards minimum energy' borrowed from the mechanical world. From this the concept of strain evolved and the requirement for a reliable quantitative basis. Experimental procedures did not appear to be able to supply the necessary data either because of limited applicability (diffraction and spectroscopic methods) or because of a distinct shortage of accurate work (thermochemical methods). On the theoretical side, quantum mechanical equations were too complex to solve in the short term and semi-empirical quantum mechanics, relying on geometries from other sources, evolved.

Molecular mechanics (MM) calculations currently provide the most accessible method. In essence the MM technique regards molecular systems as being composed of particles held together by harmonic forces. Suitable

potential functions are used to make these forces represent the different interactions occurring between the constituent atoms. The total molecular energy may then be minimised with respect to the geometry by balancing the various potentials involved.

Although the molecular mechanics method is not as sophisticated as the quantum mechanical approach, it can serve as an interim guide to correlating structure, energy, reactivity, and other molecular properties until such time as quantum mechanics is able to compete with regard to computer time. It must be stressed that MM is a model only and **Altona** (1) for instance, sees it as a natural extension to mechanical models, the short comings of which are well known. The major limitation of MM is its dependance upon empirical data.

4.2 Potential functions

The total steric energy E_s in any isolated molecule may be defined in terms of various contributors:

$$E_s = E_b + E_{nb} + E_a + E_t + E_{opbe} + E_q \quad (1)$$

where E_s = total steric energy

E_b = energy arising from bond stretching interactions.

E_{nb} = " " non-bonded "

E_a = " " angle bending "

E_t = " " torsional "

E_{opbe} = " " out of plane bending "

E_q = " " coulombic "

Each of these contributors can be represented in terms of a potential function.

(a) Bond stretching

This type of deformation has a very obvious connection with the classical Hooke's law and if it is assumed that harmonic restoring forces are in operation for small changes in bond length then,

$$E_b = \frac{1}{2}k_b (l_0 - l)^2 \quad (2)$$

where k_b = the bond stretching force constant.

l_0 = the "equilibrium" bond length for the two atoms concerned.

l = the actual bond length.

Under normal circumstances very little bond stretching is encountered and this is a consequence of the very high values for k_b (typically 200 - 1200 K.cal/mole \AA^{-2}).

Hooke's law tends to overestimate the energy required for large stretching deformations and a cubic term may be inserted to compensate. This is discussed more fully under the heading of angle bending (below).

(b) Non-bonded interactions

Numerous functions have been proposed over the years to describe non-bonded interactions because many approximations are required in order to interpret experimental results and consequently, such data are sparse. Several examples of the fit between proposed functions and observations are given in an early paper by Hendrickson (2). The data for inert gases are the easiest to interpret and

it has been the usual practice to interpolate these to cover other atoms. Thus, C..... C non-bonded interactions have been obtained, from observations of neon atoms, by curve fitting techniques.

Van der Waals produced his corrections to the gas law in 1873 by assuming that two distinct non-bonded effects were taking place. These effects were due to short range repulsive forces $\underline{E}(R)$ and a longer range attractive tendency $\underline{E}(A)$. Thus:

$$E_{nb} = E(R) - E(A) \quad (3)$$

In terms of the Lennard-Jones function this becomes:

$$E_{nb} = A/r^n - B/r^6 \quad (n = 8, 12) \quad (4)$$

where \underline{r} is the internuclear separation and \underline{A} and \underline{B} are constants.

An alternative expression was derived by Buckingham:

$$E_{nb} = A \exp(-Br) - c/r^6 \quad (5)$$

The equations (6) and (7) due to Hill (3) are preferred for practical use because they contain only two variables:

$$E_{nb} = \mathcal{E}(r^1/r)^n - 2.0 \mathcal{E}(r^1/r)^6 \quad (6)$$

$$E_{nb} = 8.28 \times 10^5 \mathcal{E} \exp(-r/0.0736r^1) - 2.25 \mathcal{E}(r^1/r)^6 \quad (7)$$

where \mathcal{E} = a force representing the ease or otherwise of moving the two non-bonded atoms.

r = interatomic distance.

r^1 = sum of the Van der Waals radii

For ($n = 12$) in equation (6) the two expressions yield curves which only differ significantly in the extreme repulsive part.

(c) Angle bending (Baeyer strain)

The basic equation is derived from Hooke's Law

$$E_a = \frac{1}{2} k_a (\theta_0 - \theta)^2 \quad (8)$$

The terms being analagous with those in equation (2).

Typical values for k_a are 0.02 K.cal per mole per degree².

Comparison with values quoted above for k_b gives an indication of the relative ease with which bond angles may be deformed.

As mentioned previously, large deformations present problems particularly for the angle bending function. Incorporation of a cubic term to equation (8) will correct the deformation in some cases. Its exact form is empirical and probably force field dependant. For instance Schleyer (4) uses the following expression:

$$E_a = \frac{1}{2} k_a (\Delta\theta^2 - k_a^1 \Delta\theta^3) \quad (9)$$

where $\Delta\theta = |\theta - \theta_0|$

k_a^1 = an additional force constant for the cubic term.

While Allinger (5) has included equation (10) in one of his force fields:

$$E_a = \frac{1}{2} k_a (\delta\theta^2 + k_a^1 \delta\theta^3) \quad (10)$$

where $\delta\theta = \theta_0 - \theta$

The changes made to Hooke's Law by these two

approaches is illustrated in Figure 4.1

Schleyer (4) has compared results from these two force fields and when large angle deformation is such that $\theta < \theta_0$, there is very little difference in the calculated angles. Boyd (6) has treated this situation as a separate system.

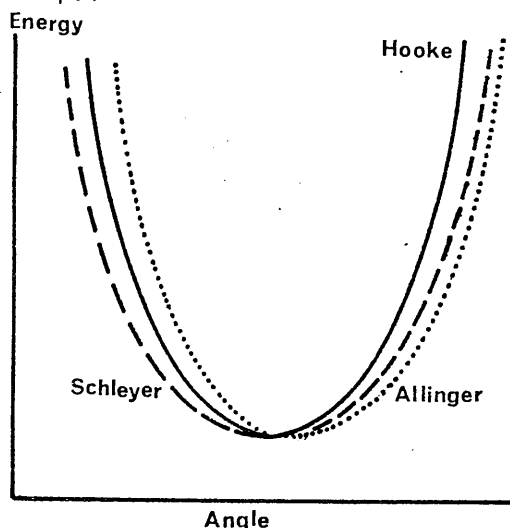


Fig. 4.1

Further difficulties arise, however, when a force field is required to cope with $\theta \gg \theta_0$ as well. Fig. 4.1 shows quite clearly that if $\Delta\theta$ is used then the angle energy contribution will decrease (equation 9) on both sides of the equilibrium position. Conversely, $\delta\theta$ (equation 10) changes in opposite senses for positive and negative deformation. Ideally, a force field should be parameterised with a wide range of compounds representing both kinds of deformation (see section 4.3).

(d) Torsion (Pitzer Strain)

It is now well known that rotation about a single carbon-carbon bond is restricted and that eclipsed and staggered conformers correspond to different energy states. The exact physical nature of such barriers is still unclear and contemporary thinking tends to favour some combination of non-bonded repulsions and electron cloud interaction. The potential function used to represent torsional interactions is:

$$E_t = \frac{1}{2}k_t (1 + \text{sign.} \cos n\omega) \quad (11)$$

where k_t = torsional force constant.

The term $(1 + \text{sign} \cdot \cos n\omega)$ represents the particular torsional situation being studied. For instance in ethane, Fig. 4.2 the staggered conformation shown has the minimum torsional interaction (zero) and this situation occurs three times during a 360° rotation. The three-fold barrier is accounted for by setting $n = 3$ in equation (11) and 'sign = +1' allows for the staggered isomer being the minimum. Hence:

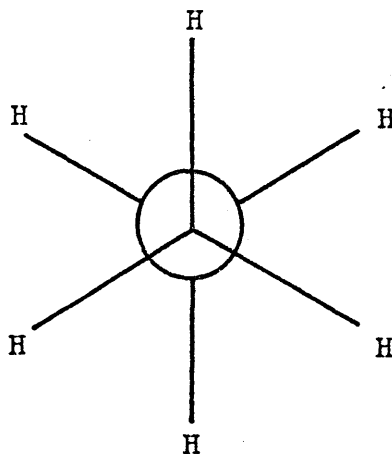


Fig. 4.2 Staggered ethane

$$E_t = \frac{1}{2} k_t (1 + 1 \cdot \cos 3\omega) \quad (13)$$

ω is the actual torsional angle which is 60° for any of the nine angles in Figure 4.2 and therefore:

$$E_t = \frac{1}{2} k_t (1 + 1 \cdot \cos (3 \times 60)) = 0$$

If an eclipsed conformation had been the minimum, as for example in propene then the three-fold barrier about the (C-C) bond

would have 'sign = -1' and represent the situation shown in Figure 4.3.

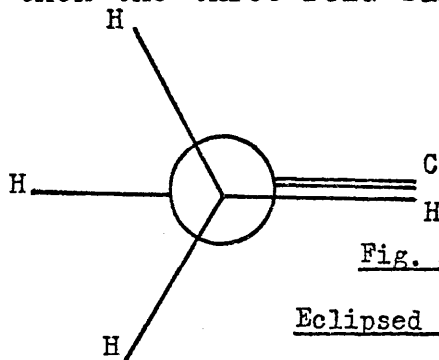


Fig. 4.3

Eclipsed propene

Equation (11) may be extended by the method of Wiberg and Boyd (7) to "drive" molecules to conformations other than the minimum so that, for example, pseudo-rotational pathways may be investigated. This is

achieved by means of equation (13).

$$E = k_t^1 \cdot \frac{1}{2} \cdot k_t (1 + \text{sign} \cdot \cos 3(\omega + \phi)) \quad (13)$$

To return to the case of ethane, discussed above, the term $(1 + \cos 3(\omega + \phi)) = 0$ for the minimum and hence:

$$\cos 3(\omega + \phi) = -1$$

giving $\omega + \phi = 60^\circ$ as one of the solutions. Substitution of suitable values of ϕ therefore gives optimised energies for reaction co-ordinates other than those of the energy minimum,

If $\phi = 0^\circ$	$\omega = 60^\circ$	equivalent to equation (11)
$\phi = 20^\circ$	$\omega = 40^\circ$	
$\phi = -20^\circ$	$\omega = 80^\circ$	

The additional force constant k^1 effectively makes the system pay a high penalty in terms of energy for any deviation from the $(\omega + \phi)$ minimum. Great caution is required in the selection of the value of k^1 . It will vary from conformation to conformation depending on, for instance, the proportion of the total steric energy contained in the torsional term concerned and on the change of energy with ω . The lowest possible value consistent with all the conformational considerations should be used. If k^1 is too large then the calculated energies will be too high or at worst numerical instability will result. The most satisfactory means for determining k^1 is by trial and error and if energy is plotted against reaction co-ordinate during the calculations then incorrect values of k^1 soon become apparent*. Equation (13) is only used to obtain the conformation required and maintain it

* See section 5.2 for further discussion

during minimisation. The actual torsional contribution to the total steric energy is calculated from equation (11).

(e) Out of plane bending

At equilibrium the three bonds about a trigonal atom such as an sp^2 carbon lie in a plane. In some instances the total strain energy in a molecule may be reduced by these bonds assuming a non-planar geometry. If the improper torsion angle χ is defined as described by Warshel, Levitt and Lifson (8) then the following function may be used:

$$E_{opbe} = \frac{1}{2} k_{opbe} (180 - \chi)^2 \quad (15)$$

where kopbe is an appropriate force constant.

(f) Coulombic interactions

Some haloketones and cyclic peptides adopt certain conformations as a result of trying to get maximum separation between like polar groups. Most cases are not so extreme as this, but nevertheless it is preferable that such interactions should be considered in calculations on molecular systems in which they are present. A potential function of the form described in equation (16) can be used:

$$E_q = (Q_1 \cdot Q_2) / (332.17 \times D \times r_{1-2}) \text{ K.cal.mole}^{-1} \quad (16)$$

where Q1, Q2 are the electrostatic charges in units of 'e' and r₁₋₂ the interatomic separation. D should strictly be the value of the dielectric constant in the region of the dipoles but the magnitude of the local quantity is not determined easily and is probably best obtained by parameterisation (see 4.3).

(g) Interdependance of functions

Clearly in a real molecule the contributors to the total steric energy cannot be treated in isolation. One of the major differences between existing applications of the MM technique is in their attitude to cross-terms. It is easy to see that if, for instance, a bond angle is changed then the non-bonded contribution from the two extreme atoms will also change and similar effects exist between other functions. From the quantum-mechanical view point, of course, the cross-terms would be taken care of automatically.

In MM the Urey-Bradley (UBFF) based force field includes all non-bonded interactions while the valence force field (VFF) algorithm ignores 1,3- and in some cases 1,4- non-bonded terms. The UBFF approach requires no cross-terms while the VFF method, to a first approximation, relies on all the cross-terms being small and accounts for the excluded interactions in the other potential functions. More sophisticated VFFs (9) have added functions for stretch-stretch and stretch-bend interactions which become significantly larger in strained environments.

4.3 Parameterisation

Once the potential functions have been selected values must be obtained for force constants and equilibrium distances and angles. The order of magnitude of many force constants is available from spectroscopic work and some distances and angles can be gleaned from crystallographic

results. Apart from this there is not a great deal of basic information at hand. This has to some extent been overcome by a trial and error method in which unknown quantities are adjusted until experimentally observed criteria (e.g. geometries, energies) are reproduced. This approach is full of pitfalls and much care is required in order to produce a balanced force field.

It is crucially important to decide what the ultimate application is to be before parameterisation is attempted. A force field (FF) which is designed to cover as many compounds as possible will not be able to give forecasts as accurate in detail as an FF based on the fewer compounds in that particular situation.

Thus to evolve an FF which can be applied in as many situations as possible, the molecules used for parameterisation should represent a balanced mixture of the different types of strain occurring.

The information which an FF is going to be required to reproduce must also be considered at this stage. Ideally the calculations should be able to obtain the physical quantities which quantum-mechanics might supply, for example:

- (a) Energies: thermochemical, strain, conformational, barriers.
- (b) Geometries: in crystals, along pathways, of different conformations.
- (c) Vibrational: frequencies, amplitudes.
- (d) Other: nmr shifts, chemical reactivity.

But again, compromise is necessary for practical purposes. Energies and geometries are probably the most important quantities from a conformational viewpoint, although attempts have been made (8) to produce consistent FFs incorporating vibrational frequencies as well.

Force field I (Appendix A) was parameterised by Miss M.J. Bovill of the Glasgow molecular mechanics group using 45 different compounds representing a balanced group of compounds for which reliable data were available. Gas phase heats of formation at 25°C were obtained by adding on group increments to convert the calculated steric energies. These increments were themselves made variables in a least squares procedure to minimise the differences between the calculated and observed heats of formation. For the 45 compounds the mean deviation in ΔH_f was 0.45 Kcal per mole.

The other FFs used in the work which follows were not so extensively parameterised but were felt to be capable of producing meaningful results. As the main interest was in the comparison of related molecules it might have been expected that the deficiencies would cancel out.

4.4 Minimisation

A parameterised FF is only capable of calculating the steric energy of a particular supplied geometry. In order to find the minimum steric energy of the isolated molecule some way must be found to allow the atomic co-ordinates to mutually adjust themselves so as to

balance the various interactions present. This may be accomplished by one of the available minimum seeking techniques.

These techniques are sometimes (10,11) referred to as 'hill climbing methods' because of their analogy to a man trying to descend a mountain on a foggy day to a specific position in a valley below. The remainder of this section is devoted to a discussion of minimisation methods.

Steepest descents

Intuitively this technique is very appealing because the minimum is sought by following the path which reduces the value of the variable being minimised. This is quite acceptable for static problems but in the dynamic multidimensional environment of MM difficulties may arise.

Steepest descents has been used by Wiberg (12) and Allinger (13-16). The energy was optimised with respect to the co-ordinates by adjusting each of the latter by some small amount in both senses and computing the resultant energy change. If an energy decrease was obtained then its value was stored and the co-ordinate returned to its unperturbed value and the process repeated until all the co-ordinates had been tested. Each atomic position was then adjusted in the direction of lower energy by an amount proportional to the partial derivative of energy with respect to the co-ordinate being moved. The cycle of testing and storing was then restarted and the whole procedure repeated until successive iterations lowered the energy by less than some prespecified amount.

The advantages of this method are that the atoms

most affecting the energy are moved the greatest amount and that 'saddle point' false minima are generally avoided. Other false minima occur, apparently fairly frequently (13,17), so that there is no absolute guarantee of reliability. Further convergence is slow, particularly if the relative weights given to the variables are not reasonably accurate and Allinger (13) has reported that the magnitude of the perturbing increment is very important with respect to the final values of torsion angles.

Pattern Search

This is very similar to steepest descents of which it may be considered a modification. The principle used is that once a valley has been found on the response surface then there is a reasonable chance that it will lead to the minimum. Steepest descents is able to locate such valleys but its ability to follow them is a function of the angle at which they are met. Pattern search was developed to surmount this problem (18).

Again co-ordinates are changed by some increment to find the direction of energy decrease but this time the disturbed co-ordinate is not restored to its original value once a lower energy position is found. The calculations then proceed to the next co-ordinate and are continued until the energy changes are sufficiently small.

Schleyer et al (19) have obtained lower energies than with steepest descents alone. Pattern search has the added advantage of being scale invariant but it can have problems locating the valleys initially and also with

following unusually shaped valleys (20). False minima are also very troublesome.

Parallel tangents

Parallel tangents is based on both the previous methods and attempts to overcome their disadvantages. From a starting point, two new locations of lower energy are sought and then the minimum on the curve through these three points is used as a new starting position. The calculation is continued until the energy changes are considered small enough.

The method has been used by Scott and Scheraga (21) and its major disadvantages are false minima and coping with curved valleys.

Non-simultaneous local energy minimisation

Allinger (22) assumed that in the area close to the minimum the potential energy surface may be represented by:

$$E = Ax^2 + By^2 + Cz^2 + Dx + Ey + Fz + G \quad (17)$$

where x, y, z are atomic cartesian co-ordinates and A to G are constants to be determined.

From a location close to the minimum another point of lower energy is found by steepest descents. The potential energy and the partial derivatives with respect to each cartesian co-ordinate are calculated and used to evaluate the constants in equation (17). If the derivative of the equation is set to zero then a new minimum position will be obtained for the atom. Successive iterations are carried out over all the atoms until the parameter shifts

are small. Allinger (22) quotes its major advantage as the saving in computer time.

Newton-Raphson Minimisation

The main difference between this and the preceding methods is that a direct solution is sought to a set of simultaneous linear equations. Iterative minimisation gives reproducible minima (1,19) and the occurrence of false minima is much less likely than with search procedures. Computer programs for calculating such direct solutions have been evolved and used by several workers (8,23,24,25). These programs have the additional advantages that quantities such as vibrational modes and thermodynamic functions may be calculated readily and that computing times are reasonable.

The MM calculations in this thesis were carried out using this technique and the following discussion is based on the program PECALC (25).

It is well known that for any continuous function $f(x)$ which has a differential coefficient at each point between $x = a$ and $x = a + h$:

$$f(a + h) = f(a) + hf^1(a) \quad (18)$$

where h is small and $f^1(a)$ is the first derivative of the function.

This is a statement of the mean value theorem and it may be extended to give the Newton-Raphson method for obtaining the roots of a polynomial by assuming that some

estimate (a) of a particular root is sufficiently close to the correct value (a + h) that equation (18) holds.

Then, since for a root: $f(a + h) = 0$:

$$0 = f(a) + hf^1(a) \quad (19)$$

In practice many iterations are needed for a complex function and if $a = x$ (trial) and $a + h = x$ (new) then equation (19) becomes:

$$x(\text{new}) = x \text{ trial} - f(a)/f^1(a) \quad (20)$$

In MM it is the energy E which must be minimised with respect to co-ordinate and hence it is the stationary value of dE/dx which is sought so that the second order form of equation (20) is required:

$$x(\text{new}) = x(\text{trial}) - f^1(a)/f^{11}(a) \quad (21)$$

where $f^{11}(a)$ is the second derivative of the function.

The situation in a real molecule is much more complex than this one - co-ordinate example since each of the constituent atoms has three co-ordinates and therefore the minimum of a multidimensional function is being sought.

Many iterations of the three dimensional analogue of equation (21) are required to obtain the minimum and if successive values of the cartesian co-ordinates are represented by x_k , x_{k+1} , then:

$$x_{k+1} = x_k - F^{-1} \nabla E(x) \quad (22)$$

and the minimum corresponds to a solution of $\nabla E(x) = 0$. The form of the matrix F in effect describes the minimisation technique in use e.g.:

$\underline{\underline{F}} = c\underline{\underline{I}}$ where $\underline{\underline{I}}$ is the identity matrix and c a properly selected constant gives the steepest descents method.

$$\underline{\underline{F}} = \frac{\partial^2 E}{\partial x_i \partial x_j} \quad i, j \leq 3 \text{ for each atom represents the three dimensional version of equation (21).} \quad (23)$$

This latter algorithm avoids the scaling problems inherent in the steepest descent approach. The calculations in Part II of this thesis have been based on equations (23). The required derivatives being calculated numerically from equation (1).

This method however, neglects co-operative movements of the atoms and may lead to errors. The energies and geometries obtained by this block diagonal calculation can be further refined by setting $\underline{\underline{F}} = \frac{\partial^2 E}{\partial x_i \partial x_j} \quad i, j = 1, 3n$ where n = number of atoms . (24)

Equation (24) thus includes all the off-diagonal terms and converges rapidly on the minimum. The full matrix technique does not converge for a crude trial structure when the components of $\nabla E(x)$ are necessarily large and the procedure adopted has been to use equation (23) initially to obtain a solution close to the stationary value and then employ the full matrix equation to calculate the minimum.

This method in addition to giving greater safeguards against false minima also avoids the difficulty encountered by some workers with torsion angle adjustment (4, 12, 26, 27). A further advantage is the facility to

calculate vibrational data from the final matrix of second derivatives.

These vibrational frequencies \underline{v}_i can be computed from their relationship to the atomic masses and potential energy. Thus, for an \underline{N} atom molecule the $(3N-6)$ vibrational frequencies are given by the eigen values of the matrix $\underline{M}^{-\frac{1}{2}} \underline{F} \underline{M}^{-\frac{1}{2}}$ where \underline{M} is the diagonal matrix of atomic masses.

Equations (25) and (26) may then be used to calculate the vibrational free energy, \underline{F}_{vib} , and the vibrational zero point energy \underline{E}_{vib}^0 respectively.

$$\underline{F}_{vib} = kT \sum_{i=1}^{3N-6} \ln \left[1 - \exp(-h\nu_i/kT) \right] \quad (25)$$

$$\underline{E}_{vib}^0 = \frac{1}{2}h \sum_{i=1}^{3N-6} \nu_i \quad (26)$$

4.5 References

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CHAPTER FIVE

Conformational analysis of a series of cyclo-octadienes

5.1 Forward

Carbocyclic systems are generally classified into groups of compounds based on the number of atoms forming the ring. The 1885 strain theory of Baeyer attempted to account for the variation in properties of cycloalkanes on the basis of the internal angles required for a series of regular polygons. This approach assumed that the cycloparaffins were planar and predicted increasing strain from (and including) 6-membered rings upwards.

The thermodynamic data that we now have available (Fig. 5.1) shows that small rings (3- to 5-membered) and medium rings (7- to 11- membered) are the most strained. Baeyer's theory only accounts for the progression observed in the small rings and it is not until allowance is made for the possibility of puckering that the heats of combustion of the larger rings can be explained.

One of the puckered forms of cyclohexane has a heat of combustion of 157.4 K.cal per mole per $-\text{CH}_2-$ group identical with the value for a straight chain alkane and this figure is also approached for the cyclic compounds with twelve or more atoms in a single ring. These large rings also closely resemble open chain paraffins. The anomalous heats of combustion of the medium ring carbocycles arise because such compounds cannot completely relieve the torsional strain by changing bond angles. In addition, there is usually some strain arising from trans-

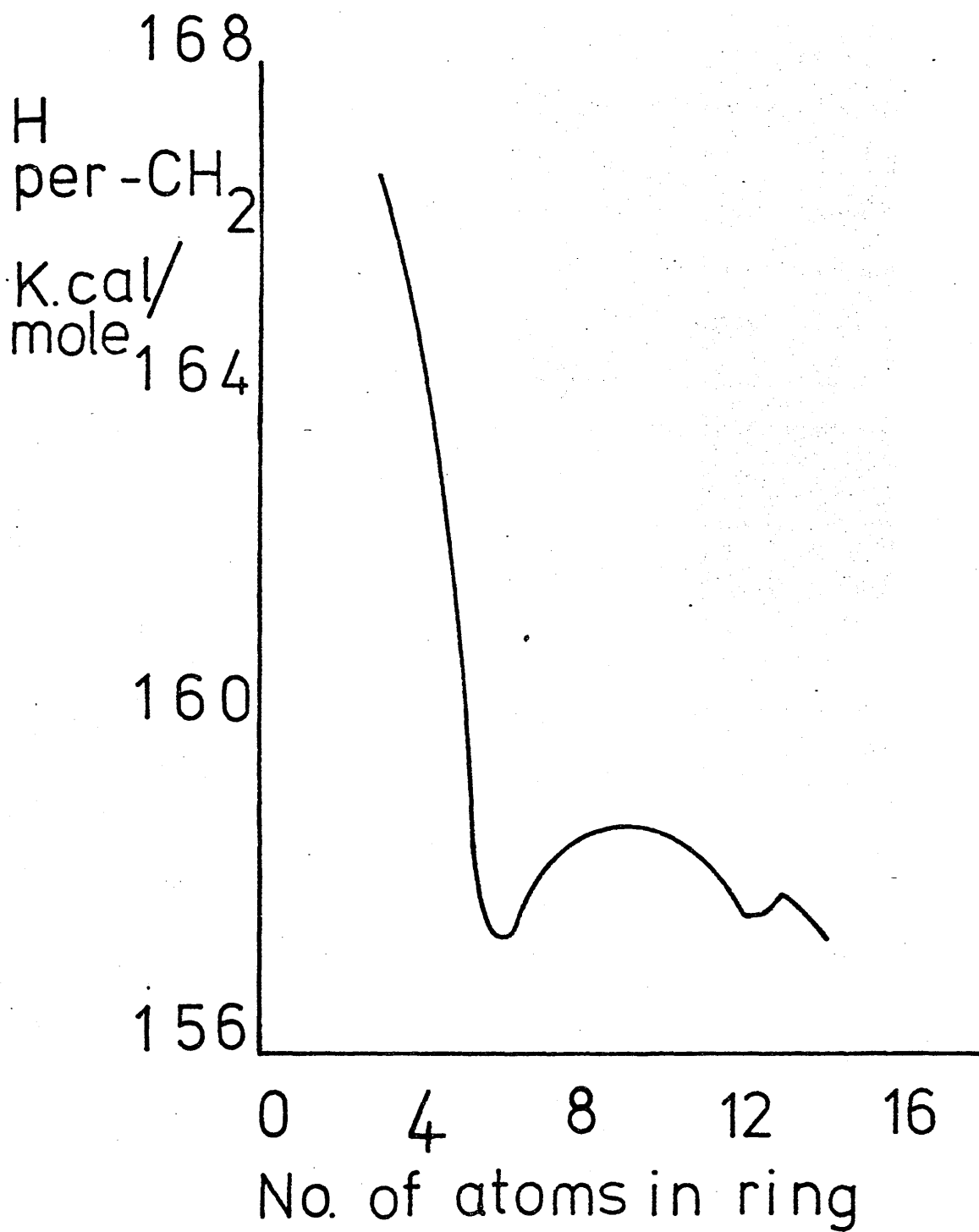


Fig. 5.1 Heats of combustion (H) of some cycloalkanes

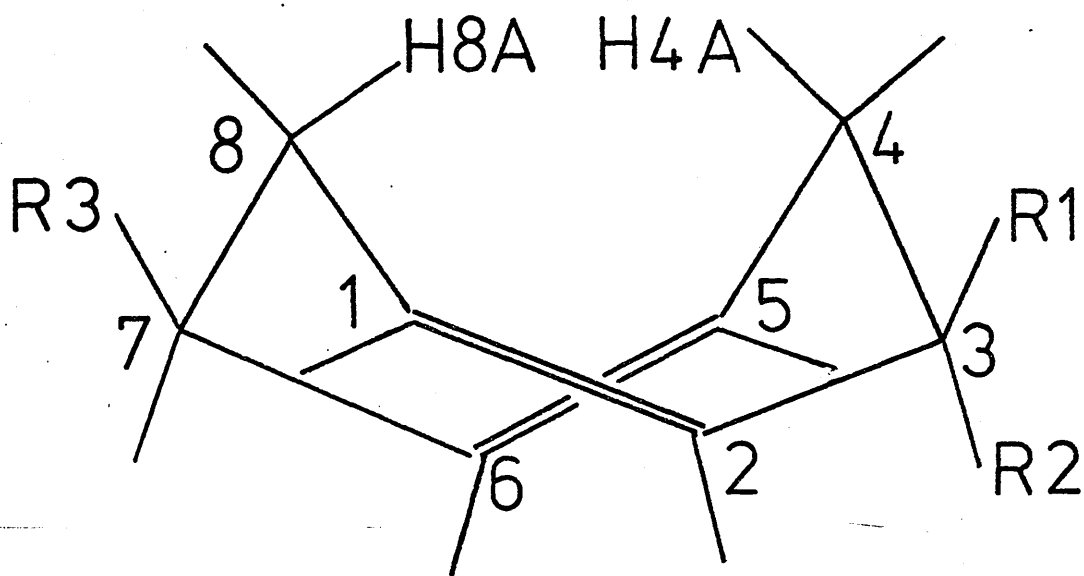
annular interactions.

Molecular mechanics (MM) by simulating the balance between these competing forces should be able to provide quantitative explanations of why one conformation is preferred over another in specific circumstances. The series of compounds studied in this chapter was selected because a previous investigation into a dibromo-octadiene had posed a number of interesting questions concerning the conformational processes which might occur in such a system. Also, the reported crystal structure of this particular compound provided sufficient data with which to check the reliability of the system to be used for subsequent calculations.

5.2 Introduction

The conformations of cis, cis-cyclo-octa-1,5-dienes have been investigated by a number of workers. Dunitz and Waser (1) used geometric arguments to show that both a rigid chair-like form and a family of flexible boats (Fig. 5.2) are possible for these dienes. In 1950 Roberts (2) made approximate calculations of the steric energies of the transition states between the boat, skew, and chair conformers of 1,6-dichloro-cyclo-octadiene. He suggested, on the basis of dipole moment measurements, that the boat was the most stable form.

The parent diene cis, cis-cyclo-octa-1,5-diene (COD) has been studied in the gas phase by electron diffraction



COD

$R_1=R_2=R_3=H$

SDBCOD

$R_1=R_3=Br, R_2=H$

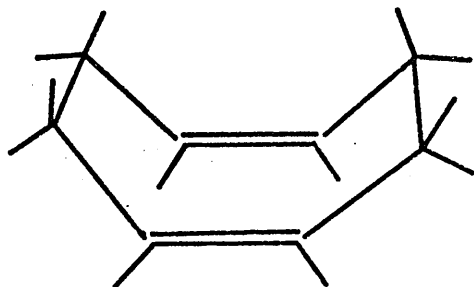
ADBCOD

$R_1=H, R_2=R_3=Br$

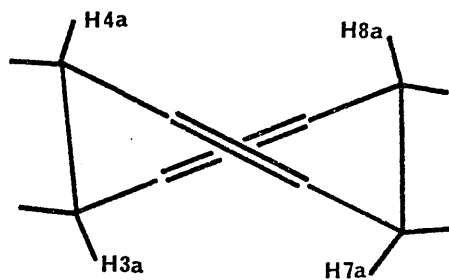
ADMCOD

$R_1=H, R_2=R_3=Me$

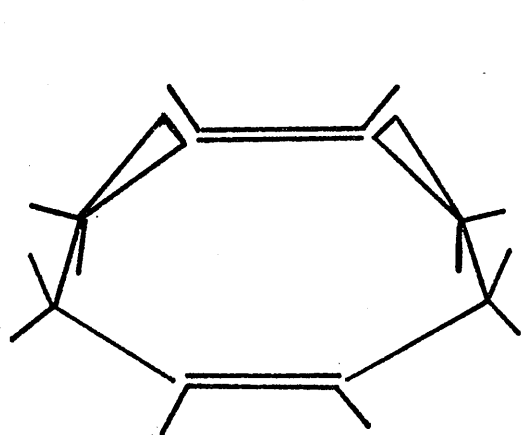
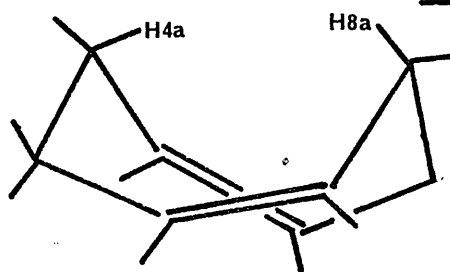
Boat \underline{C}_{2v}



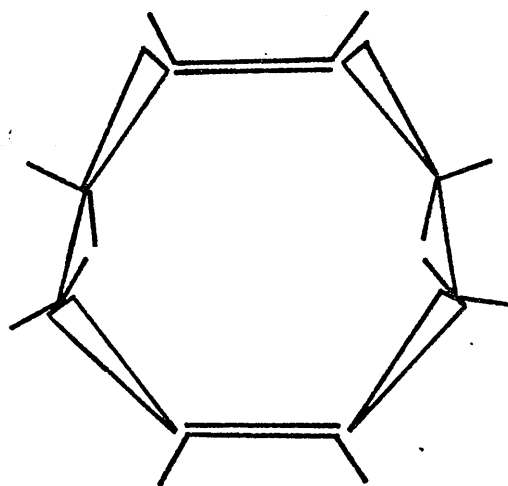
Skew \underline{D}_2



Twist-boat \underline{C}_2



Transition state
 \underline{C}_s



Chair \underline{C}_{2h}

and this indicated that an equilibrium mixture of twist boat (most predominant) and chair conformers was present (3). Anet and Kozerski (4) determined the heights of the barriers to interconversion of the stereoisomers of COD by the variable temperature nmr technique and showed that the twist boat was the minimum energy form. Pauncz and Ginsburg (5) calculated that from consideration of only the H....H non-bonded interactions the boat of COD should be preferred to the chair. MM calculations (6,7,8) have suggested that the chair is more stable than the boat and that the twist boat represents the energy minimum. A number of x-ray analyses on metal complexes of COD have also shown a preference for the C_2 boat although this does not necessarily imply that the uncomplexed cyclo-octadiene should exist in the same conformation.

X-ray studies of (all ax-) -2,6-dimethylcyclo-octa-3,7-cis, cis-diene-1,5-dicarboxamide (15) and syn-3,7-dibromo-cis, cis-cyclo-octa-1,5-diene (SDBCOD)(16) found that both these derivatives of COD exist as twist boats in the crystalline state. Other physical techniques (nmr and ir spectroscopy and dipole moment measurement) have also been applied to SDBCOD in the solid and/or liquid phases (16). The results of these investigations have all indicated that the twist boat is the major conformer present in the solid and liquid states.

Dibenzocyclo-octa-1,5-diene (DBZCOD) has also been studied by x-ray diffraction and the eight-membered ring was found to have adopted the chair conformation (17). Recently, strain energy calculations (9, 18) have been

used to interpret the nmr spectrum of DBZCOD (19,20,21). The calculations explained the nmr spectra of both COD and DBZCOD in terms of an interconversion process between isoenergetic twist boats. The routes by which this can occur are discussed more fully in section 5.5. The differences between the preferred interconversion pathways of these two compounds were attributed to the distinct torsional barriers about the $C_{sp^2} - C_{sp^3}$ bonds of the octadiene ring.

The preparation of anti-3,7-dimethyl-cis,cis-cyclo-octa-1,5-diene (ADMCOD) has been reported (22) and the conformational interconversion processes are currently being investigated by variable temperature nmr spectroscopy (23) using a sample kindly supplied by Professor Heimbach (Gesamthochschule Essen and Max-Planck-Institut für Kohlenforschung).

The strain energy calculations in the literature (e.g. 7, 8, 18) have thus considered only the parent diene and derivatives such as DBZCOD which do not represent all the possible conformational situations arising during twist boat interconversion. In the work which follows the MM technique has been extended to include these other conformations for 3,7-disubstituted-derivatives of COD and the results compared with the available experimental data.

The latest Glasgow alkane/alkene FF(FFI) (24) was used for COD and ADCOD while the less extensively parameterised FFII (24) was employed for SDBCOD and ADBCOD since FFI did not include any terms for interactions

involving the bromine atoms. Pseudorotation was achieved by the method of Wiberg and Boyd (25) and plots of energy against reaction co-ordinate (the C(2) - C(3) - C(4) - C(5) torsion angle) were used to check for false minima as described in Chapter 4. This approach does not necessarily give the true minimum energy for transition states because, even though there may be some regular change in energy as different constraints are imposed on the model, full relaxation is not possible. Ermer (26) has recently pointed out that when the torsion angle used as reaction co-ordinate is symmetrically located in relation to a two-fold axis of symmetry (2AS) in the nearest minimum structure, and this same symmetry element is not present in the transition state, then the method of Wiberg and Boyd will impose this 2AS on that transition state. One of the examples he cites in connection with this is the boat-chair transition state of cyclohexane. Wiberg and Boyd (25) used their "driving method" to calculate energies for the two transition states corresponding to \underline{C}_2 and \underline{C}_s symmetry by assuming that four- and five- atoms were respectively coplanar and, (with a single C-C-C-C torsion angle as reaction co-ordinate)⁺ found that the former was preferred by about 0.5 K.cal mole⁻¹. Pickett and Strauss (27) on the other hand derived a pseudorotating transition state which incorporated both \underline{C}_2 and \underline{C}_s forms implying that the energy difference between them was much smaller than that reported by Wiberg and Boyd. Ermer repeated the work of Wiberg and Boyd and obtained energies for the two transition states using the technique of removing the constraints at an appropriate mapping point. He obtained geometries which were close to, but not exactly corresponding to, the rigid four- and five- atom

⁺ for the \underline{C}_2 TS

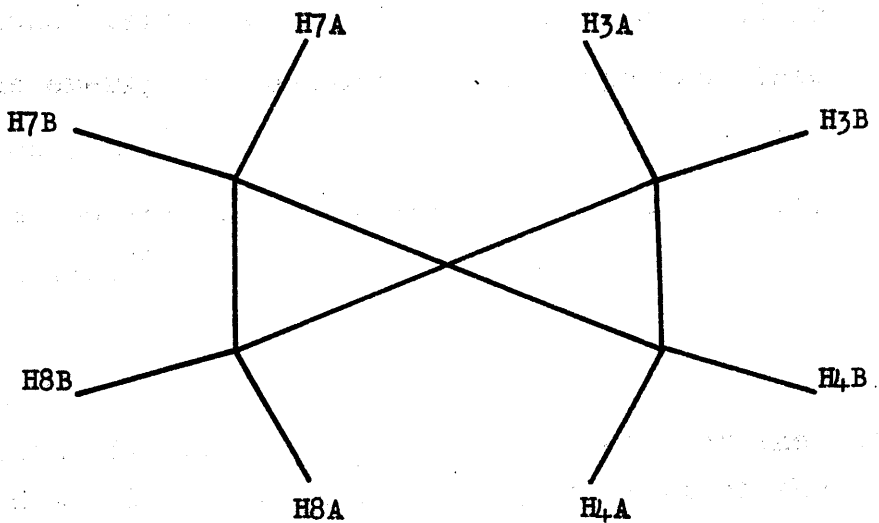
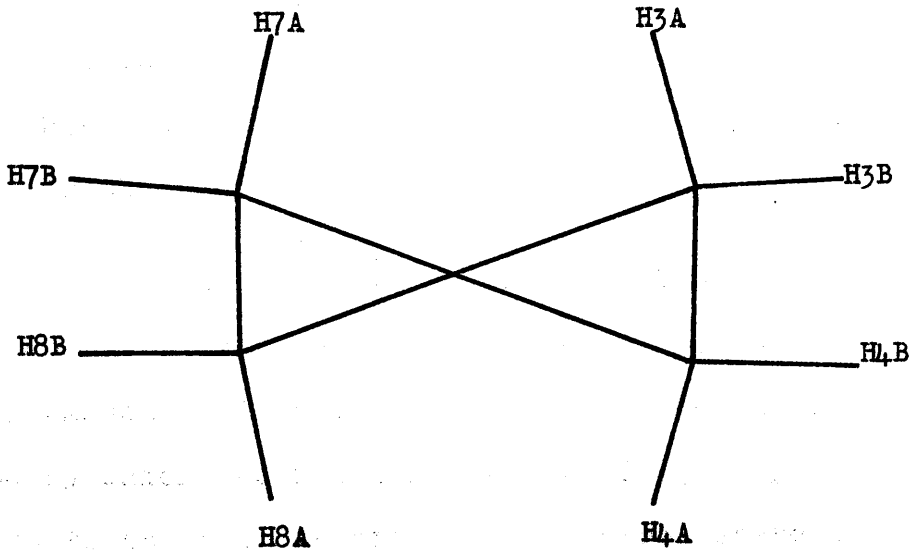
coplanar structures. Moreover the energy difference was now 0.00086 K.cal in favour of the C_s structure.

This result clearly indicates that caution is required when using mapping techniques to calculate conformational barriers. The calculations on cyclo-octadienes reported in this thesis constrained torsion angles which were symmetrically located about the C_2 axis of the global minimum and the possibility therefore arises that the calculated transition states do not represent the true maximisations of the potential energy function, especially as the derivatives were calculated numerically. Therefore as a final check the constraints were removed from the models for the boat, skew, and boat-to-chair transition states of COD and ADMCOD. There was a reduction in energy in every instance, but in particular, the skew forms displayed drastic changes in both energy and geometry. The constrained (C) and non-constrained (NC) skew conformations of COD for example have an energy difference of over 6 K.cal per mole and this is sufficient to change the skew from being the least- to being the most-preferred intermediate between twist-boats*.

Ermer also discusses the relationship between the C_{2v} boat (B) and C_2 twist-boat of COD with respect to the calculations published by Allinger and Sprague (8, 28) and comments that

*The original calculations had constrained all nine torsion angles about the C(3)-C(4) and C(7)-C(8) bonds, thus 'locking' the conformation of the substituent hydrogens. It was found for COD that constraining the carbon skeleton only (e.g. C(2)-C(3)-C(4)-C(5)) gave essentially the same result as was obtained on subsequent removal of all the constraints. It appears that the fully constrained model (i.e. all nine torsion angles) is equivalent to Allinger's method of fixing the coordinates and has the same dire consequences.

(NC)



(G)

they initially described the former as a minimum (with respect to the C_{2h} chair; they did not consider the C_2 twist boat) but correctly reported the B conformation as a transition state between twist boats in the later paper (8). However, in the same report they state that the boat is more stable than the skew, a result now shown to be incorrect for the reasons outlined above.

In some of the calculations which follow the variation in steric energy with reaction co-ordinate for the fully constrained model is illustrated and it is now clear that the method by which the results were obtained only allows conclusions to be drawn regarding the general shape of the energy versus torsion angle curves. Accordingly, the calculations for the dibromo-octadienes are only completely reliable as far as the geometries and energies of the non-constrained twist boats are concerned. In cases where the conformational barriers are of prime interest (COD and ADMCOD) the energy and geometry of the important intermediates have been obtained by using the mapping technique to approach the maximum and then removing the constraints to obtain full relaxation.**

**On occasions it was found that the geometry of the fully constrained model was not close enough to that of the transition state for convergence. In such cases the constraints were removed from the eight exocyclic torsion angles about the C(3) - C(4) and C(7) - C(8) bonds and a new stationary point sought with the block diagonal algorithm. This new 'trial' geometry was then sufficiently close to the maximum for convergence to occur for the completely unconstrained system under full matrix conditions.

5.3 syn- 3,7- Dibromo - cis, cis-cyclo-octa-1,5 - diene
(SDBCOD)

Method of calculation

The program PECALC (29) was used to minimize the steric energy in FFII (24) with respect to the atomic co-ordinates (16). The mathematical model was pseudorotated by the method of Wiberg and Boyd (25) and the minimum energy calculated at 10° intervals throughout the pseudorotational pathway. Initially the block diagonal algorithm was employed for minimisation. Subsequently, the energies and geometries of important stereoisomers on possible interconversion routes were verified and refined by the full matrix calculation described in Section 4.4. The constraints were not removed during the final stages of refinement.

Results and Discussion

The study of SDBCOD by Mackenzie et al (16) reported the global minimum of this compound as being TB1 (Fig. 5.3). This has been confirmed by MM calculations and the geometry of this conformation is compared with the x-ray analysis in Table I* and serves as an indicator of the reliability of FFII.

*The geometric parameters cannot be compared rigorously because of errors inherent in the x-ray results ($R = 11.1\%$). The possibility of differences between the calculated isolated molecule and the observed crystal structure must also be considered in this context.

Fig. 5.3 SDECOD: Variation of energy with conformation



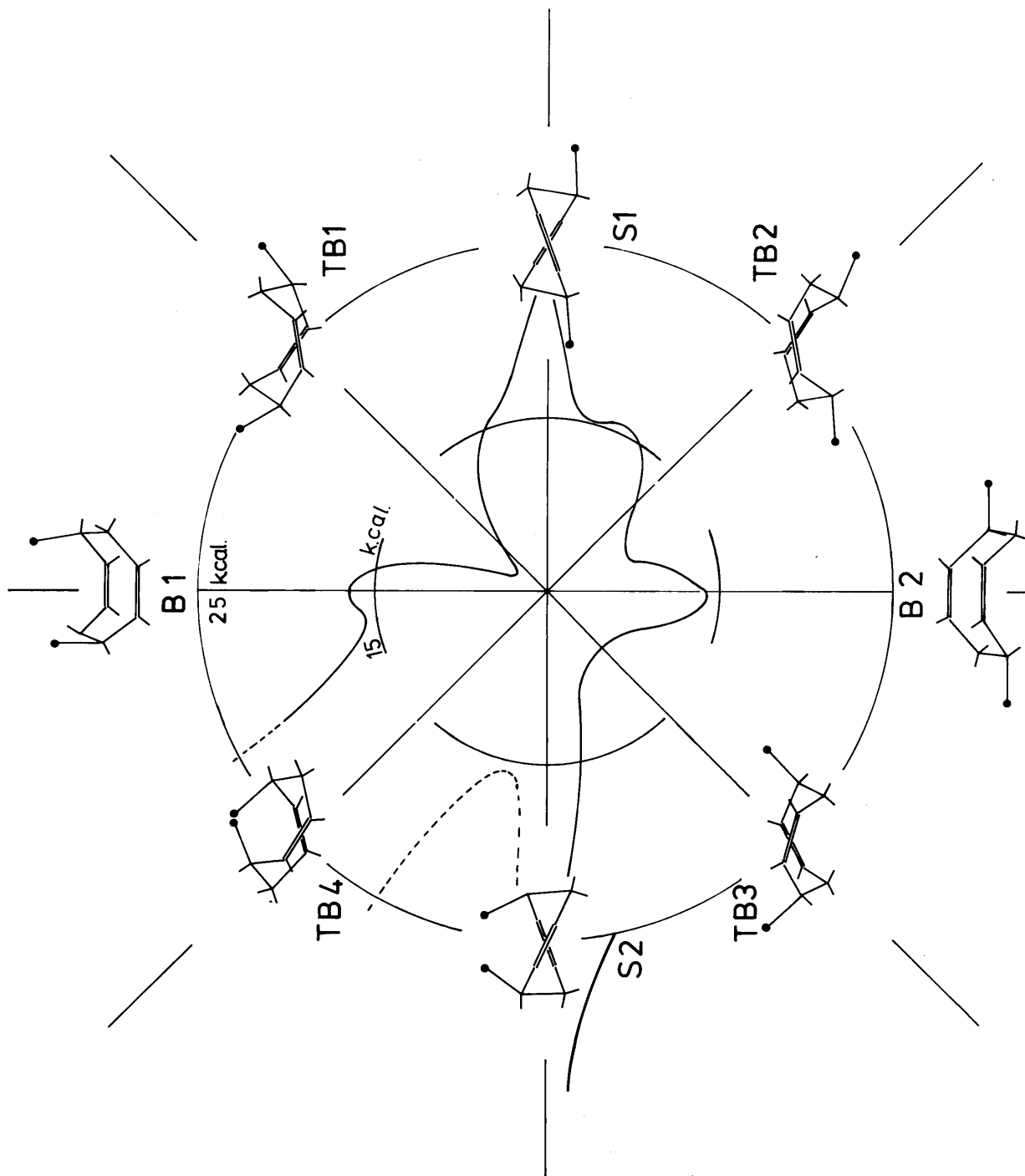


TABLE I SDBCOD: COMPARISON OF OBSERVED AND CALCULATED STRUCTURES

PARAMETER	CRYSTAL STRUCT (16)	CALCULATION
BOND LENGTHS (A)		
C(1) - C(2)	1.37(3) *	1.336
C(2) - C(3)	1.48(3) **	1.504 **
C(3) - C(4)	1.51(3)	1.543
BOND ANGLES (DEG.)		
C(1) - C(2) - C(3)	129(2) **	126.1 **
C(2) - C(3) - C(4)	116(2) **	113.5 **
NON-BONDED DISTS (A)		
C(4)...C(8)	3.04 +	3.03
TORSION ANGLES (DEG.)		
C(2)-C(3)-C(4)-C(5)	65 ++	42
* ESDS IN PARENTHESIS ** AVERAGE VALUE + FROM IR SPECTRUM ++ NMR IN SOLUTION ALSO GAVE 65 DEGREES		

Fig. 5.3 shows a radial plot of the variation of steric energy with the C(2) - C(3) - C(4) - C(5) torsion angle. The curve has been interpolated (broken line) where necessitated by the inability of the calculation to produce a minimum when severe van der Waals forces are present (e.g. TB4). For reasons noted earlier the relative energies between conformations should be treated with extreme caution particularly in the vicinity of the skew conformations where full relaxation produces a considerable reduction in strain energy by allowing the close-contact transannular atoms (e.g. H3A and H7A for NC) to move further apart. Nevertheless, the results do indicate the existence of three local minima (TB1, TB2 and TB3) which could interconvert and thus support the observation of Mackenzie et al (16) that in solution, at least, more than one twist boat is present. Fig. 5.4 illustrates some of the more probable interconversion pathways and Table II gives the relative energies of the more important intermediates together with the calculated barriers to interconversion.

In view of the uncertainty of the calculations for the boat, skew and TSA and TSB conformations (all of which were not allowed to fully relax) it would be improper to comment further on these conformational barriers.

It is apparent however that the chair form is less stable than twist boats TB1, TB2 and TB3.

1. TB1 \rightarrow TSA \rightarrow CHAIR \rightarrow TSB \rightarrow TB2

1A. TB1 \rightarrow TSA \rightarrow CHAIR \rightarrow TSB \rightarrow TB3

2. TB1 \rightarrow B1 \rightarrow TB4

3. TB1 \rightarrow S1 \rightarrow TB2

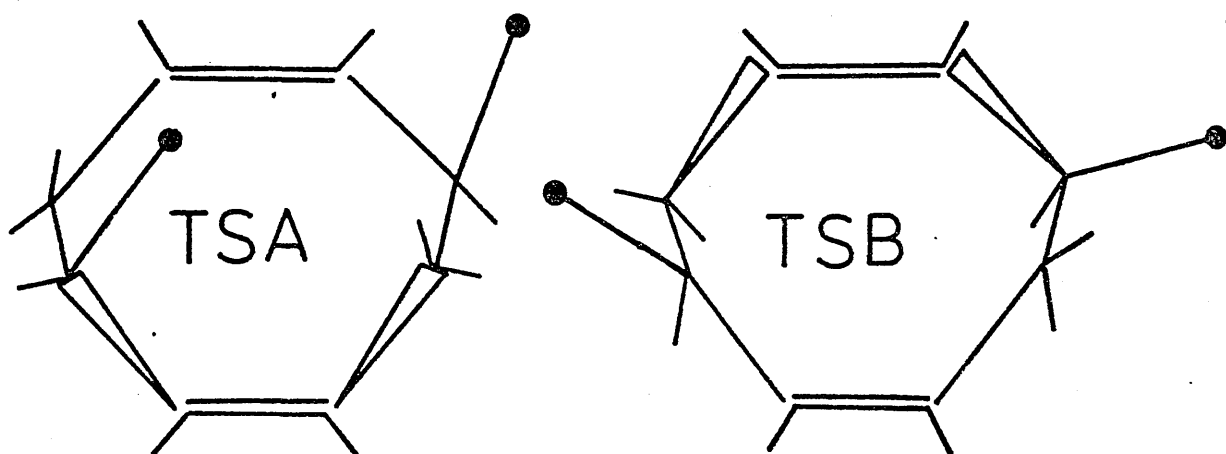


Fig. 5.4 SDECOD: Pathways for Twist-boat Interconversion.

TABLE II SDCOD: RELATIVE ENERGIES

CONFORMER	REL. ENERGY *	ENERGIES IN K.CAL/MOLE
TB1	0.00	
TB3	2.41	
TB2	3.05	
CHAIR	4.90	
TSA	5.21	
TSB	6.83	
B2	7.97	
B1	10.14	
S1	18.63	
S2	39.25	

*

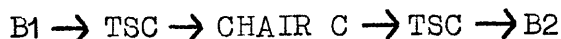
The calculations show that there are two local
minima (TB1 and TB3) and that the diaxial con-
former is more stable than the diequatorial variety. This is

5.4 anti-3,7-Dibromo-cis, cis-cyclo-octa-1,5-diene (ADECOD)

Results and Discussion

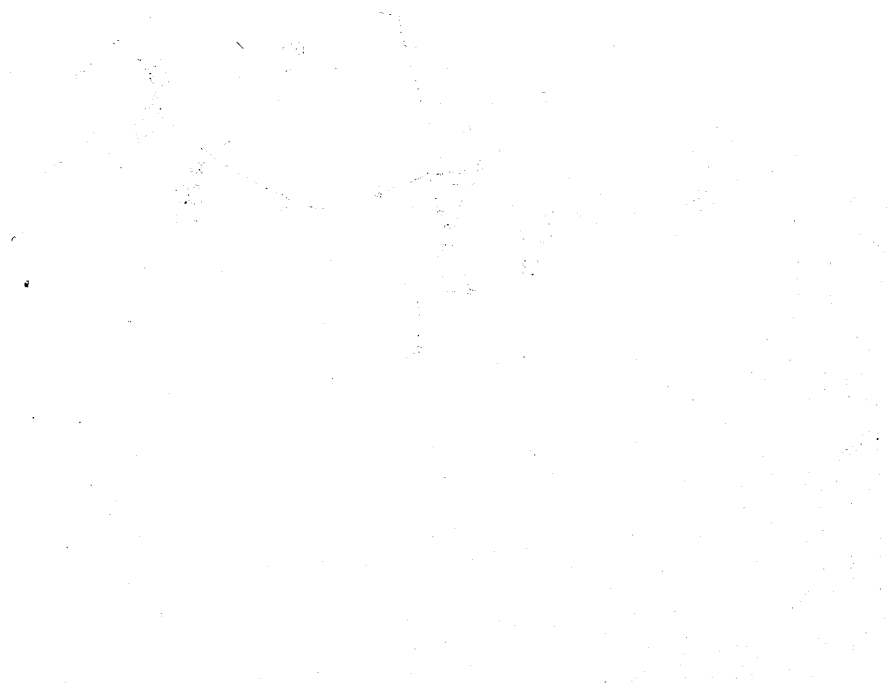
It has, to date, not been possible to isolate this compound (23). Nevertheless, in anticipation of ADECOD being available at some future date it was decided to investigate its conformational behaviour. MM calculations were carried out as described for SDECOD and the results are shown in Fig. 5.5 and Table III. All the remarks made with reference to the possibility of only partial maximisation having been achieved for certain conformations of SDECOD apply equally to ADECOD.

Anti - as opposed to syn - substitution of the bromine atoms at C(3) and C(7) has no symmetry and this results in there being two possible chairs (C and D) in which the bromine atoms can be either diaxial (C) or diequatorial (D). The corresponding boat-to-chair transition states (also called C and D for convenience) either move the bromine atoms closer together (C) or further apart (D) from their relative positions in the boat. Inspection of models indicates that only the following unique conformational changes can occur through the chairs:



The calculations show that there are two isoenergetic global minima (TB1 and TB3) and that the diaxial chair is more stable than the diequatorial variety. This latter result is unexpected and appears to be connected with the

Fig. 5.5 ADECOD: Variation of energy with conformation



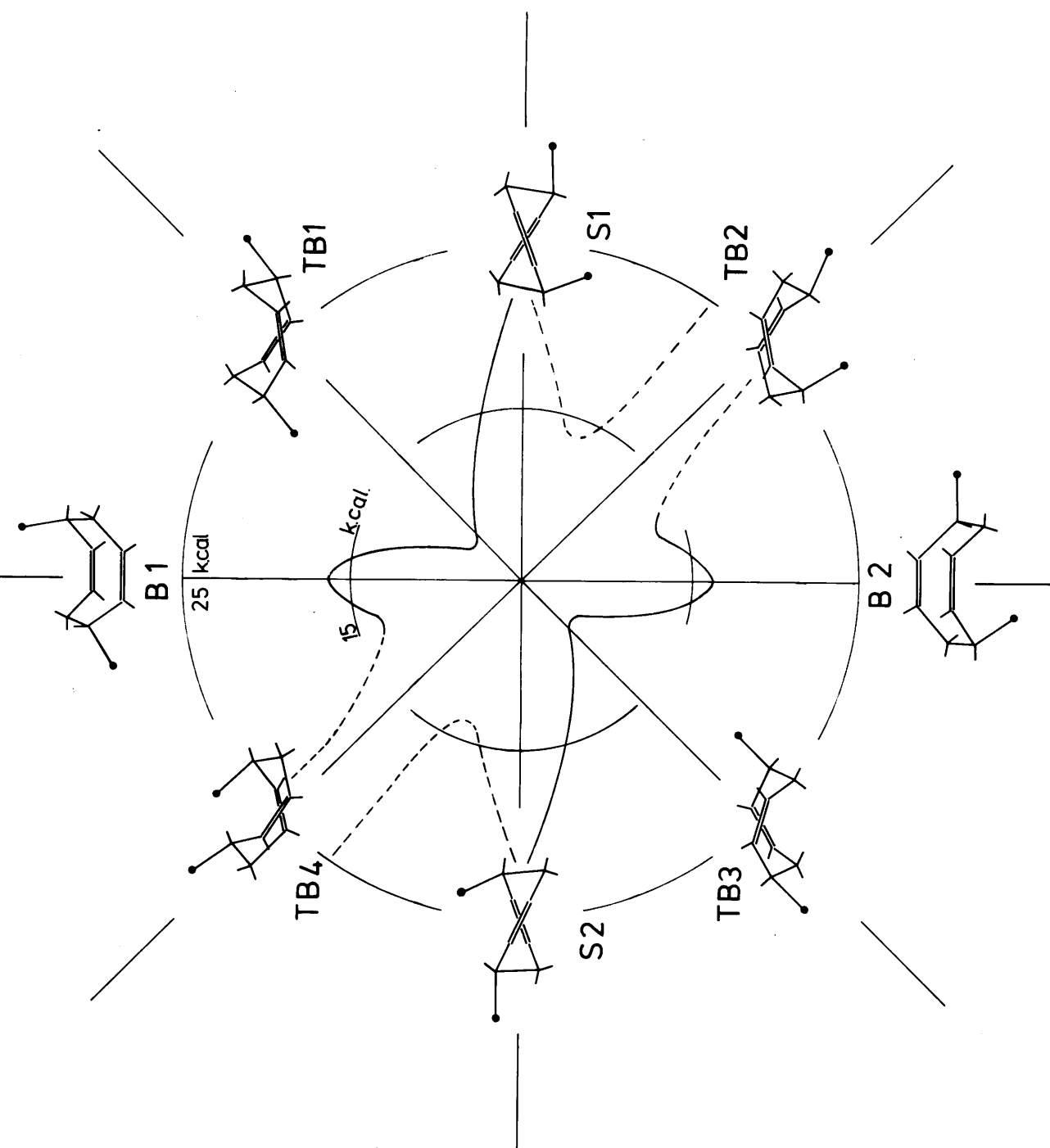


TABLE III ADBCOD: RELATIVE ENERGIES

CONFORMER	REL. ENERGY *
TB1	0.00
CHAIR C	3.34
CHAIR D	4.20
TSC	3.80
TSD	5.49
BOAT	8.53
SKEW	26.42

ENERGIES IN K.CAL/MOLE

*

CHAIR
C

CHAIR
D

1. TB1 \rightarrow TSC \rightarrow CHAIR C \rightarrow TSC \rightarrow TB3

1A. TB1 \rightarrow TSD \rightarrow CHAIR D \rightarrow TSD \rightarrow TB3

2. TB1 \rightarrow B1 \rightarrow TB4

3. TB1 \rightarrow S1 \rightarrow TB2

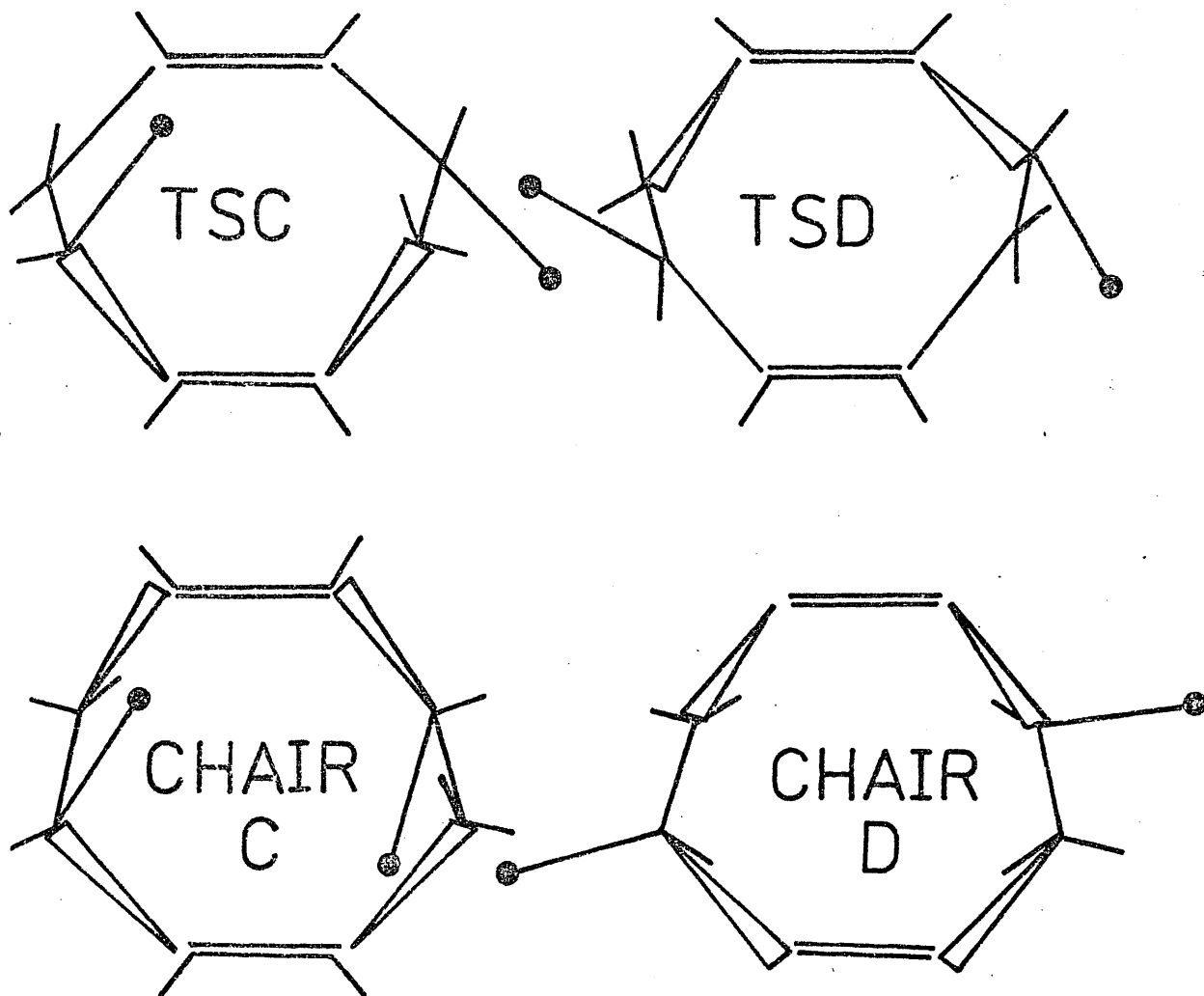


Fig. 5.6 ADECOD: Pathways for Twist-boat Interconversion.

large van der Waals radius of bromine which in chair D is responsible for substantial non-bonded strain between the bromine atoms at C(3) and C(7) and one of the substituents on the α - carbon atoms.

Once again it is not possible to comment on the barriers for the routes between twist boats (Fig. 5.6).

The relative energies listed in Table III (and Table II) for the constrained boat, skew and TSC and TSD, would almost certainly be reduced under fully relaxed conditions although it is not possible to estimate the relative reduction, and hence the relative stabilities, of each conformer under these conditions.

5.5 cis, cis-Cyclo-octa-1,5-diene (COD)

Introduction

The calculations reported in sections 5.2 and 5.3 served as a preliminary test of the reliability of the FF and facilitated the installation of the suite of computer programs. It was decided to apply this system to the parent hydrocarbon (COD) because both an electron diffraction study (3) and nmr measurements of conformational barriers (4) in COD were available as further tests. Initially FF II was employed to map the pseudorotational circuit and calculate energy differences between key conformations. Subsequently the more extensively parameterised Glasgow alkane/alkene force field (FFI) was used and it is these later calculations that are reported here.

Results and Discussion

The calculated conformation of the twist boat is compared with the geometry observed in the electron diffraction study in Table IV. The agreement is good and the value of the C(2) - C(3) - C(4) - C(5) torsion angle (52°) is close to that of 45° measured in the gas phase (3).^{*} The x-ray analysis of SDBCOD (16) reported the corresponding angle as 65° . Allinger and Sprague (8) and Favini et al (7) have reported angles of 38° and 27° respectively and Ermer, using the Lifson-Ermer FF calculated a value of 55° (30). The uncertainty regarding the electron diffraction result prohibits the use of this angle as an unambiguous test for comparison of the various FFs but the results from FFI and the Lifson-Ermer FF lie somewhere between 45° and 65° and although the connection between an isolated molecule of COD and its crystal syn-3,7-dibromo derivative is tenuous the correlation of the calculated geometry of COD with the electron diffraction results provides evidence of the reliability of FFI.

The calculations support the view that the twist boat geometry represents the steric energy minimum. In the rigid chair, for instance, the torsional strain can only be relieved by an increase in Baeyer strain but in the C_{2v} boat the eclipsing around the C(3) - C(4) and C(7) - C(8) bonds can be reduced by pseudorotation. The decrease in energy arising from the twisting around these bonds is offset by the increasing approach of H_{4a} and H_{8a} (for TB1) and the conformation adopted by the minimum energy boat is directly

* See note (a) Table IV for comments regarding the accuracy of this torsion angle.

TABLE IV COD: COMPARISON OF OBSERVED AND CALCULATED STRUCTURES

PARAMETER	ELECTRON DIFFN(3)	CALCULATION
BOND LENGTHS (A)		
C(1) - C(2)	1.341	1.340
C(2) - C(3)	1.502 *	1.512 *
C(3) - C(4)	1.554	1.530
BOND ANGLES (DEG.)		
C(1) - C(2) - C(3)	127.8 *	127.8 *
C(2) - C(3) - C(4)	116.8 *	115.8 *
NON-BONDED DISTS (A)		
C(1)...C(5)	3.302	3.39
C(1)...C(6)	3.047	3.08
C(2)...C(6)	3.318	3.31
C(3)...C(7)	3.815	3.81
C(4)...C(7)	3.191	3.19
C(4)...C(8)	3.131	3.05
H(4)...H(8)	----	1.94
TORSION ANGLES (DEG.)		
C(2)-C(3)-C(4)-C(5)	45	52

* AVERAGE VALUE

a CALCULATED FROM THE GEOMETRY QUOTED IN REF.3. ASSUMES THAT VALENCE ANGLES C(2)-C(3)-C(4) AND C(3)-C(4)-C(5) ARE EQUAL. THIS IS KNOWN TO BE INCORRECT FROM THE CRYSTAL STRUCTURE OF SDBCOD(16) AND FROM THE CALCULATIONS. THE ELECTRON DIFFRACTION RESULTS THAT ARE AVAILABLE GIVE ONLY AN AVERAGE VALUE FOR THESE ANGLES. HENCE THE ACCURACY OF THE C(2)-C(3)-C(4)-C(5) TORSION ANGLE IS NOT KNOWN.

related to the H.....H non-bonded potentials used in the FF(32). The figures in Table IV indicate that this strain can be relieved by deformation of the ring angles at the sp^2 and sp^3 carbons to 127.8 and 116.8 respectively. The two hydrogens are then moved further apart (calculated separation 1.94\AA) while the C(4) and C(8) transannular distance is increased from ca. 2.3\AA , apparent from an undistorted Dreiding model, to 3.131\AA reported in the electron diffraction study. The calculated value is 3.05\AA and the **x-ray analysis** of SDBCOD(16) gave this distance as 3.04\AA .

The balance achieved in the minimum energy boat is changed if the pseudorotation is continued towards the skew form since there is an increase in the Pitzer strain around C(3) - C(4) and C(7) - C(8), and in the skew form itself there are two pairs of hydrogens (H4a, H8a; and H3a, H7a) in close contact. The need for extreme caution when calculating the strain energy of intermediates such as the skew has been discussed from the symmetry view point in section 5.2. In purely mechanical terms, full relaxation permits co-operative movements of the atoms and the illustrations of the geometry of the non-constrained (NC) and fully constrained (C) skew conformations of COD show that this results in a reduction of the H...H transannular repulsions. Inspection of the calculated structures indicates that this is achieved by a flattening of the ring.*

The calculation has thus been very successful in

*The energy calculated for the constrained molecule is correct for that molecule but this structure is not a major intermediate on the pseudorotational pathway because a substantial reduction in strain energy is achieved by adoption of the NC geometry.

reproducing the geometry of the twist boat and for COD there are four of these (isoenergetic) twist boats on the pseudorotational circuit. Only one quadrant of this circuit is unique but because each twist boat has a different pair of methylene hydrogens in close contact it is possible, by use of the low temperature nmr technique, to observe and measure any interconversion of twist boats which might be taking place. This is precisely what has been done by Anet and Kozerski (4) and their spectra are shown in Fig. 5.7. The ^{13}C nmr spectra for the methylene carbon resonance changes from a single peak at high temperatures to a 1:1 doublet at -176° . The symmetries of the boat, chair, and skew forms are such that methylene carbons are always equivalent and therefore the doublet at -178° must represent a twist boat (the methylene carbons in the boat to chair 6 co-planar transition state are also non-equivalent but because this is more unstable than either the boat or the chair it would not be observed as a single entity). The singlet at high temperatures is therefore consistent with rapid interconversion of twist boats (process A, with $\Delta G = 4.2 \pm 0.2$ K. cal mole $^{-1}$ at -176°) resulting in each of the methylene carbon pairs (C(3), C(7) and C(4), C(8)) interchanging steric environments.

The possible pathways for such interconversion are given in Figure 5.8 and the pmr spectra of the methylene protons show that two of these conversion processes are occurring simultaneously (at room temperature), one with a

$\Delta G = 4.4 \pm 0.1$ K. cal mole $^{-1}$ at -177° (process B) and the other with $\Delta G = 4.9 \pm 0.1$ K. cal mole $^{-1}$ at -168° (process C).

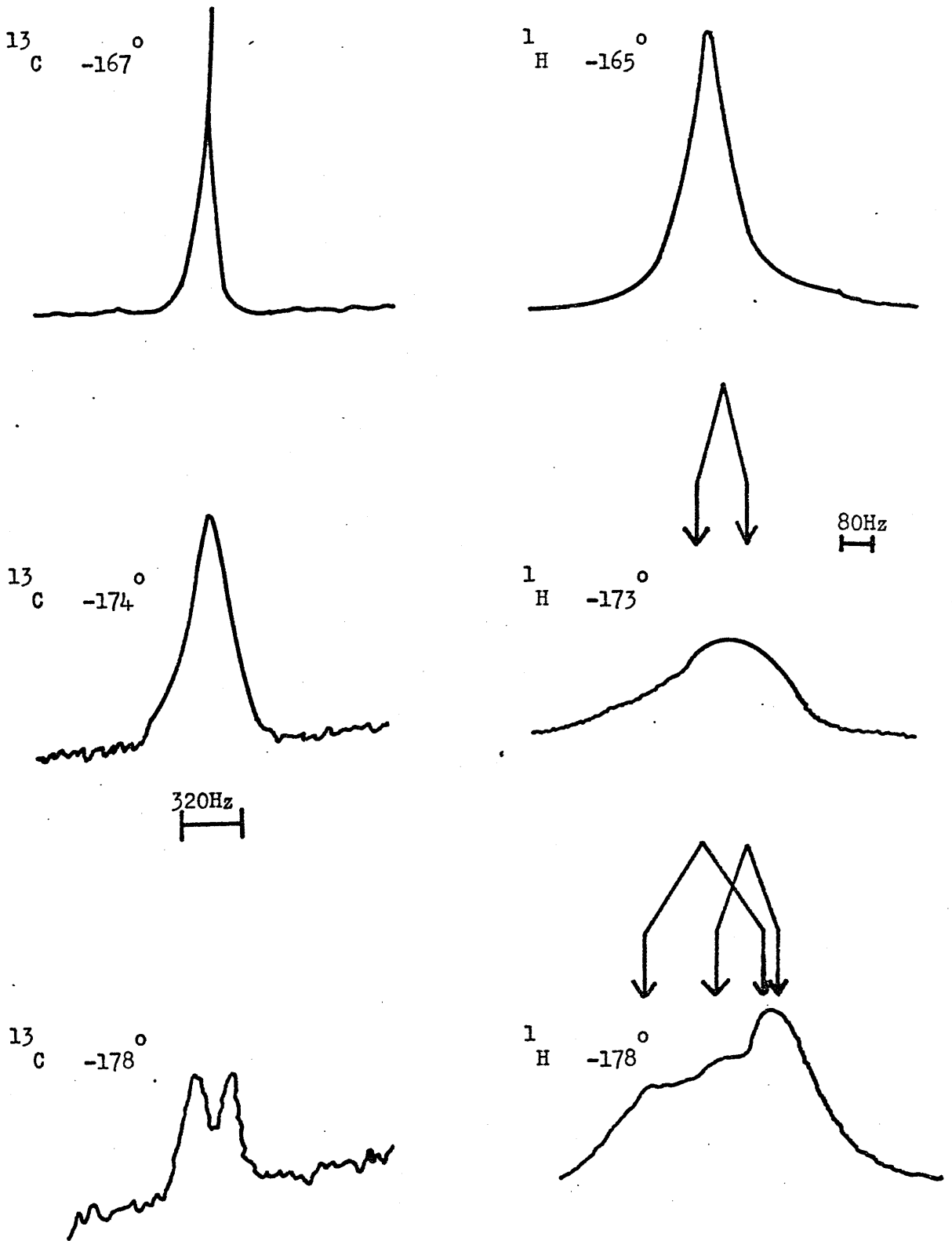
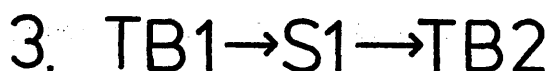
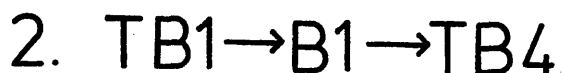
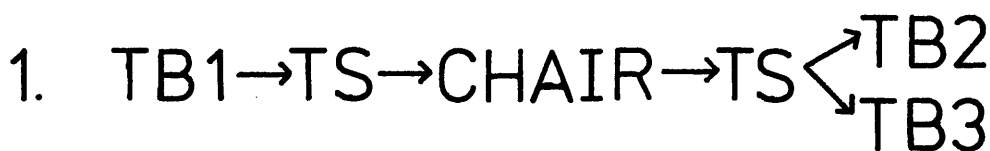


Fig. 5.7 COD: Variable Temperature NMR Spectra of Methylene Groups(4)



Since the components have different shapes, they do not represent route 1 since there is no symmetry (the one formed being not 3).
 Approximate equilibrium time is significant in the comparison of the results of complete averaging.

Two other possibilities therefore exist:

Process 1 represents Route 2

and

Process 3 represents Route 1 or Route

OR

Process 1 represents Route 1

and

Process 3 represents Route 2 or Route 1

Fig. 5.8. COD: Pathways for Twist-boat Interconversion.

Imagine the number processes 1 and 2. In the paper, Alst and Hoxerski state that the path between the processes 2 and 1 can in principle be determined. The processes remain stable

The assignment* of the three spectral processes:

A	$\Delta G = 4.2 \text{ K cal mole}^{-1}$ at -176°C
B	$\Delta G = 4.4 \text{ K cal mole}^{-1}$ at -177°C
C	$\Delta G = 4.9 \text{ K cal mole}^{-1}$ at -168°C

to specific modes of twist boat interconversion:

1	$\text{TB1} \rightarrow \text{B1} \rightarrow \text{CHAIR} \rightarrow \text{B2} \rightarrow \text{TB}$
2	$\text{TB1} \rightarrow \text{B1} \rightarrow \text{TB4}$
3	$\text{TB1} \rightarrow \text{S1} \rightarrow \text{TB2}$

is not unambiguous. However, the ^1H spectrum at -177°C (process B) indicates that the two components have different chemical shifts and hence cannot represent route 1 since there are two possible chairs (the one formed being determined by which group of 6 atoms is coplanar in the transition state) leading to complete averaging.

The following possibilities therefore exist:

Process B represents Route 2

and

Process C represents Route 1 or Route 3

OR

Process B represents Route 3

and

Process C represents Route 1 or Route 2

Process A observed in the ^{13}C spectrum must involve the sum of processes B and C. In their report Anet and Kozerski state that the path representing process 2 "can in principle be determined ... by accurate strain energy

*e.g. Process A is not necessarily assigned to route 1.

calculations".

These workers have now carried out such calculations and their results, together with those from other sources (7, 8, 30), are shown in Table V. There is broad agreement between the predictions of Ermer (30), Anet and Kozerski, and this work regarding the relative order of the conformational barriers. The calculations of Favini et al (7) were only concerned with computing the energies of various isolated conformers of COD and therefore cannot be applied to this discussion. Allinger and Sprague (8) find the skew as the most unstable conformation; an identical result to that obtained by the author for the fully constrained model, thus suggesting that they have not allowed co-operative movements of the atoms.

The calculations reported in this thesis indicate that:

Process B	represents route 3 (skew)
Process C	represents route 2 (boat)*

As noted previously there are discrepancies between the observed and calculated barriers and these are of about the same order as those found by other workers.

In the following discussion the nmr spectra are explained in terms of these assignments.

*The energy difference between routes 1 and 2 is only 0.1 K.cal and therefore this assignment is not unambiguous. However, both Ermer, and Anet and Kozerski also find route 2 to be preferred to route 1 (and Ermer obtains a difference of 0.2 K.cal between them, in the same sense).

TABLE V COD: RELATIVE ENERGIES FROM VARIOUS SOURCES

CONFORMER	RELATIVE ENERGY *			THIS WORK
	FAVINI(7)	ALLINGER(8)	ANET(31)	ERMER(30)
TB	0.00	0.00	0.00	0.00
CHAIR	2.75	1.45	---	0.77
BOAT	3.84	3.05	---	4.55
SKEW	6.44	5.94	---	2.27
TRANSITION STATE	---	5.23	---	4.64
ROUTE 1	---	5.23	7.2	4.64
ROUTE 2	3.84	3.05	6.6	4.55
ROUTE 3	6.44	5.94	4.2	2.27

MEASURED BARRIERS BY NMR(4)

	BARRIER	POSSIBLE COMBINATIONS OF ROUTES		
		ROUTE 2	ROUTE 3	ROUTE 1 OR ROUTE 2
PROCESS B	4.4 *	ROUTE 1 OR ROUTE 3	ROUTE 2	ROUTE 1 OR ROUTE 2
PROCESS C	4.9 *			

* ENERGIES IN K.CAL/MOLE

The ^{13}C spectrum results from the simultaneous occurrence of (1) TB1 to TB2 conversion (Fig. 5.7) through the skew and (2) TB1 to TB4 via the boat taking place less frequently.* This spectrum of the methylene carbons gives only their interchange in environment resulting from twist boat interconversion since the symmetries of the intermediate skew and boat prevent their conformations being observed in ^{13}C spectroscopy. The pmr spectrum, however, distinguishes between the two pathways. The methylene proton band at -165°C is composed of a single peak because at that temperature it is possible to interconvert between all four twist boats and so on average each of the eight methylene protons can occupy all the possible steric environments. The calculations have indicated that lowering the temperature to -173°C prevents the route 2 process from taking place. This produces the doublet observed since four of the methylene protons are always "outside" (in the sense that they cannot come into close contact as H4a and H8a in TB1) while the other four are always "inside". Thus this rocking motion via the skew between TB1 and TB2 which is still taking place at -173°C averages the shifts of the "inside" protons and also averages the "outside protons" giving the observed spectrum. The activation energy for this process is too high to be overcome at -178°C and the ^1H spectrum at this temperature is given by the four different types of methylene proton present in an isolated twist boat.

*Strictly, all three processes are occurring but for simplicity the discussion has been restricted to the two major interconversions.

5.6 anti-3,7-Dimethyl-cyclo-octa-1,5-diene (ADMCOD)

Introduction

The results for COD show that FFI is able to reproduce the energies and geometries of its various conformational possibilities. As stated previously, FFI cannot at present be used for strain energy calculations on SDBCOD and ADBCOD because it has not been parameterised for the bromine interactions. The lack of experimental data for ADBCOD (see section 5.4) prevented the predictions for that compound being tested. However, the recent synthesis of ADMCOD (22) and the co-operation of Professor Heimbach in supplying a sample for nmr analysis provided an opportunity for further study of both the reliability of FFI and of the conformational situation in anti-3,7-derivatives of COD.

Results and discussion

The full nmr results are not yet available because of equipment problems but preliminary spectra indicate that there is just one main interconversion process occurring with a barrier of about 5 K.cal per mole (23).

Some of the possible processes are shown in Fig. 5.9 and the calculated relative energies of the major conformers are listed in Table VI. The conformational variety possible in ADMCOD is exactly as described previously for ADBCOD. Thus the relationship of the transition states and chairs to each other and to the diequatorial or diaxial location of the bromine atoms of ADBCOD remains unaltered on the anti- substitution of the methyl groups.

In ADMCOD utilisation of the full matrix relaxation technique has shown that TB₂, TB₄, the boat (of approximate C_{2v} symmetry), and the skew are all local stationary points (either maxima or inflexions) and the present calculations give little indication of the form of the energy vs. reaction coordinate curve between them. Fortunately route 1A through the diequatorial chair (D) represents the minimum energy route

1. TB1 \rightarrow TSC \rightarrow CHAIR C \rightarrow TSC \rightarrow TB3

1A. TB1 \rightarrow TSD \rightarrow CHAIR D \rightarrow TSD \rightarrow TB3

2. TB1 \rightarrow B1 \rightarrow TB4 \rightarrow S2 \rightarrow TB3

3. TB1 \rightarrow S1 \rightarrow TB2 \rightarrow B2 \rightarrow TB3

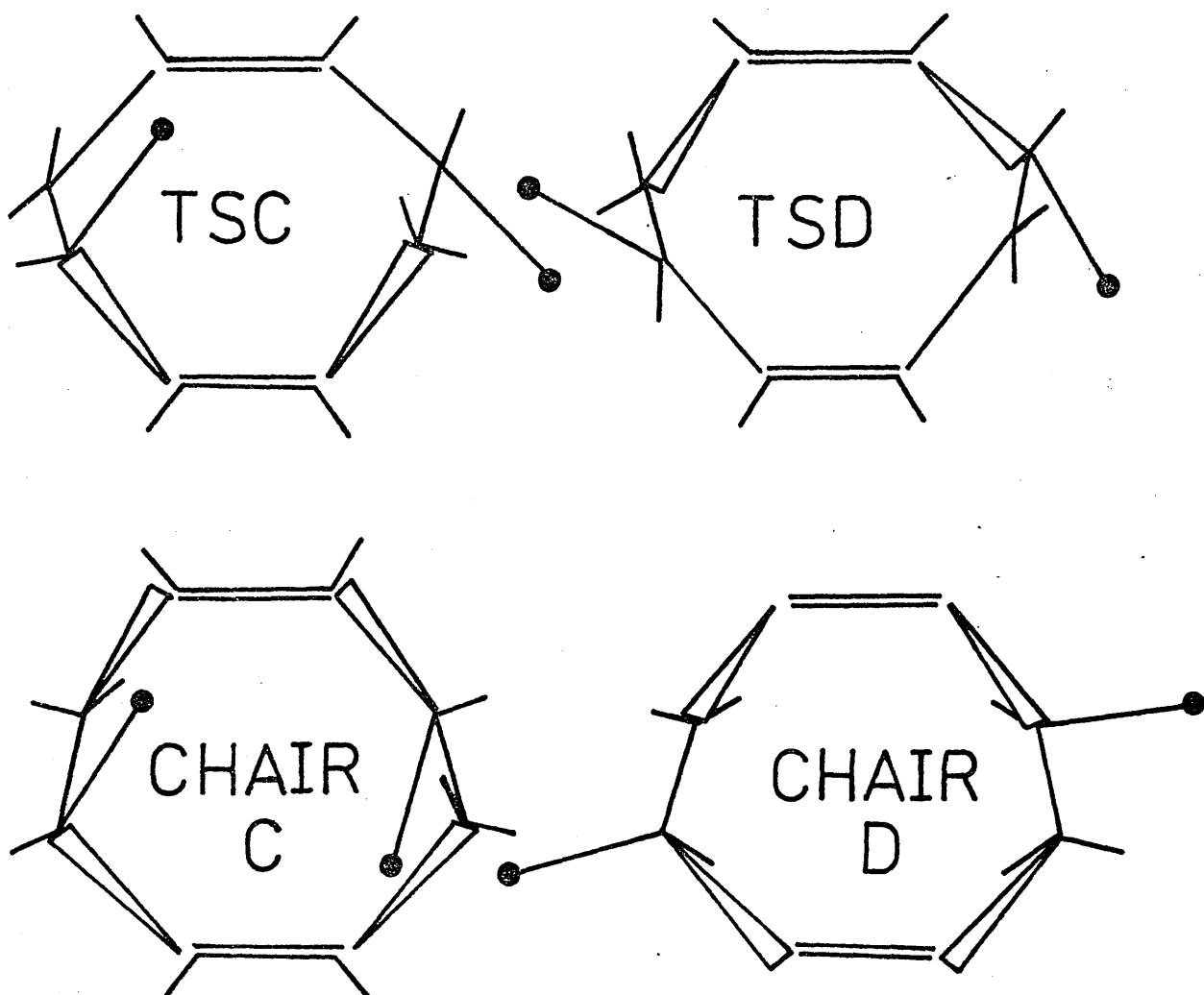


Fig. 5.9 ADMCOD: Pathways for twist-boat interconversion

TABLE VI ADMCOD: RELATIVE ENERGIES

CONFORMER	REL. ENERGY *	
TB1	0.00	
CHAIR D	0.95	
CHAIR C	2.26	
TSD	4.98	
TSC	9.02	
TB2/TB4	6.61	
BOAT	5.78	
SKEW	13.03	
ROUTE 1	9.02	(THROUGH CHAIR C)
ROUTE 1A	4.98	(THROUGH CHAIR D)
ROUTE 2	6.61	(THROUGH BOAT AND TB4)
ROUTE 3	13.03	(THROUGH SKEW AND TB2)
		ENERGIES IN K.CAL/MOLE

*

As shown in Figure 1, the relative energies of the conformers are given in kcal/mole. The energy of the chair conformation is taken as zero. The energy of the boat conformation is 5.78 kcal/mole. The energy of the skew conformation is 13.03 kcal/mole. The energy of the chair conformation is 0.00 kcal/mole. The energy of the chair conformation is 0.95 kcal/mole. The energy of the chair conformation is 2.26 kcal/mole. The energy of the chair conformation is 4.98 kcal/mole. The energy of the chair conformation is 9.02 kcal/mole. The energy of the chair conformation is 6.61 kcal/mole. The energy of the chair conformation is 5.78 kcal/mole. The energy of the chair conformation is 13.03 kcal/mole.

and this is therefore indicated as the main process observed in the low temperature nmr. It is not clear however whether the approximate C_{2v} boat is a necessary precursor of the chair*.

Inspection of Dreiding models has suggested that the 6-coplanar transition state** of Roberts can be formed from a member of the boat family in which the C(2)-C(3)-C(4)-C(5) torsion angle is 10 or even 20° and more evidence is required before this anomaly can be resolved. It would be particularly interesting to have some information on the processes taking place (and the barriers to them) in more sterically crowded 3,7-anti-substituted octadienes which should have a larger energy differential between the boat and TSD.

* Anet (ref. 30 p571) has remarked that the lowest barrier for an interconversion process might only be reached in some cases by driving several torsion angles simultaneously and unsymmetrically.

**See section 5.7 for further discussion of the exact conformation of the transition state.

5.7 Comparison of results

The employment of two different FFs for the calculations on COD and its 3,7 disubstituted derivatives restricts the degree to which comparisons can be made. The improvements in the more recent of the two FFs (FFI) may be expected to affect the proportion of the total strain arising from each type of contributor rather than the relative order of the various conformers. The initial calculations for COD and ADMCOD were carried out with FFII so that results are available from both FFs for these two compounds and reinforce the previous statement. The discussion which follows is nevertheless not detailed because it is based on calculations performed with both FFI (COD and ADMCOD) and FFII (SDBCOD and ADBCOD).

The minimum energy conformation is the twist boat for all four compounds although the value of the C(2) - C(3) - C(4) - C(5) torsion angle varies between 42° and 52° . This is not altogether unexpected since this angle is dependant on the magnitude of H... H interaction in the FF used (32) and also the precise effects of (i) the differences between the environments in which the observed values of 45° and 65° were measured and (ii) the substitution in the series, are not known.

The calculated C(4) C(8) separations and the endocyclic angle deformation are consistent with the experimental geometries (3, 16).

Substitution of the anti methyl groups at C(3) and

C(7) has little effect on the relative stabilities of most of the conformers encountered. It does, however, change the major intermediate for twist boat interconversion from the skew (COD) to the diequatorial chair (ADMCOD) because of the strain imposed on the TB2 and TB4 forms of ADMCOD.

Both the boat intermediate and TB4 are, in the case of ADMCOD, distorted conformations of the 'purer' forms obtained for COD, and this is a reflection of the extra strain imposed on the molecule by the methyl substitution.

The boat to chair transition state proposed by Roberts does not have six atoms in one plane for either COD or ADMCOD. Unconstrained full matrix minimisation has indicated that these six atoms, C(3) to C(8), have a maximum of four atoms lying in one plane, the other two atoms being displaced slightly from this plane. This situation is very similar to that suggested by Ermer (26) for cyclohexane.

The relative stabilities of the chair conformations of ADBCOD and ADMCOD requires comment. It was noted in Section 5.4 that the diaxial chair was preferred for ADBCOD, and this was explained on the basis of the non-bonded strain between the bromines and one of the α -hydrogen atoms in the diequatorial chair. In ADMCOD this latter chair is more stable since the methyl substitution changes the contact distances and can also reduce these interactions by rotation about the C(3) - and C(7)-methyl bonds. The diaxial chair of ADMCOD has more van der Waals strain than

the corresponding conformer of ADBCOD as a result of the increased number of additional atoms ($-\text{CH}_3$ as opposed to $-\text{Br}$) and their greater proximity to the octadiene ring.

The general agreement between the various FFs in interpreting the spectra for COD is encouraging, although the calculated barriers are not as close to the measured values as might be desired. Nevertheless, with the exception of Allinger's calculations,* the other independent studies of COD (26,31) find the same relative order of preference for the three pathways as reported here and enable some confidence to be placed in the predictions made by FFI for ADMCOD.

*The reasons for this have been discussed in Sections 5.2 and 5.5.

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CHAPTER SIX

Studies of germacranolide stereochemistry

6.1 Introduction

The considerable body of experimental data available on the stereochemistry of sesquiterpenes (e.g. 1,2,3) has revealed several phenomena of which only a small number have been interpreted.

The work reported in this chapter was carried out in order to investigate certain of the observed peculiarities of germacranolides. In particular some aspects of the stereochemistry of those germacranolides possessing an α -methylene (or methyl) - γ - lactone fused to a trans, trans,-cyclodeca-1,5 diene ring (Fig. 6.1) have been put on a quantitative basis by the application of molecular mechanics (MM).

The macrocycle in these systems is known to occur frequently in two of the possible conformations of trans, trans,-cyclodeca-1,5,diene (I). Conformation A (Fig. 6.2) has been observed in, for example, costunolide (4), elephantol (5) and alatolide (6) while conformation B is typified by dihydromikanolide (7) and shiromodiol (8) but, apart from some preliminary calculations by Favini (9), very little is known about the relative stabilities of the various conformations of I. In addition the trans, trans-double bonds are deformed by consistently different amounts from the ideal value of 180° for the C—C=C—C torsion angles (10) and the data currently available are insufficient to show whether this is caused by the substitution of the lactone ring and/or the other substituents to I, or

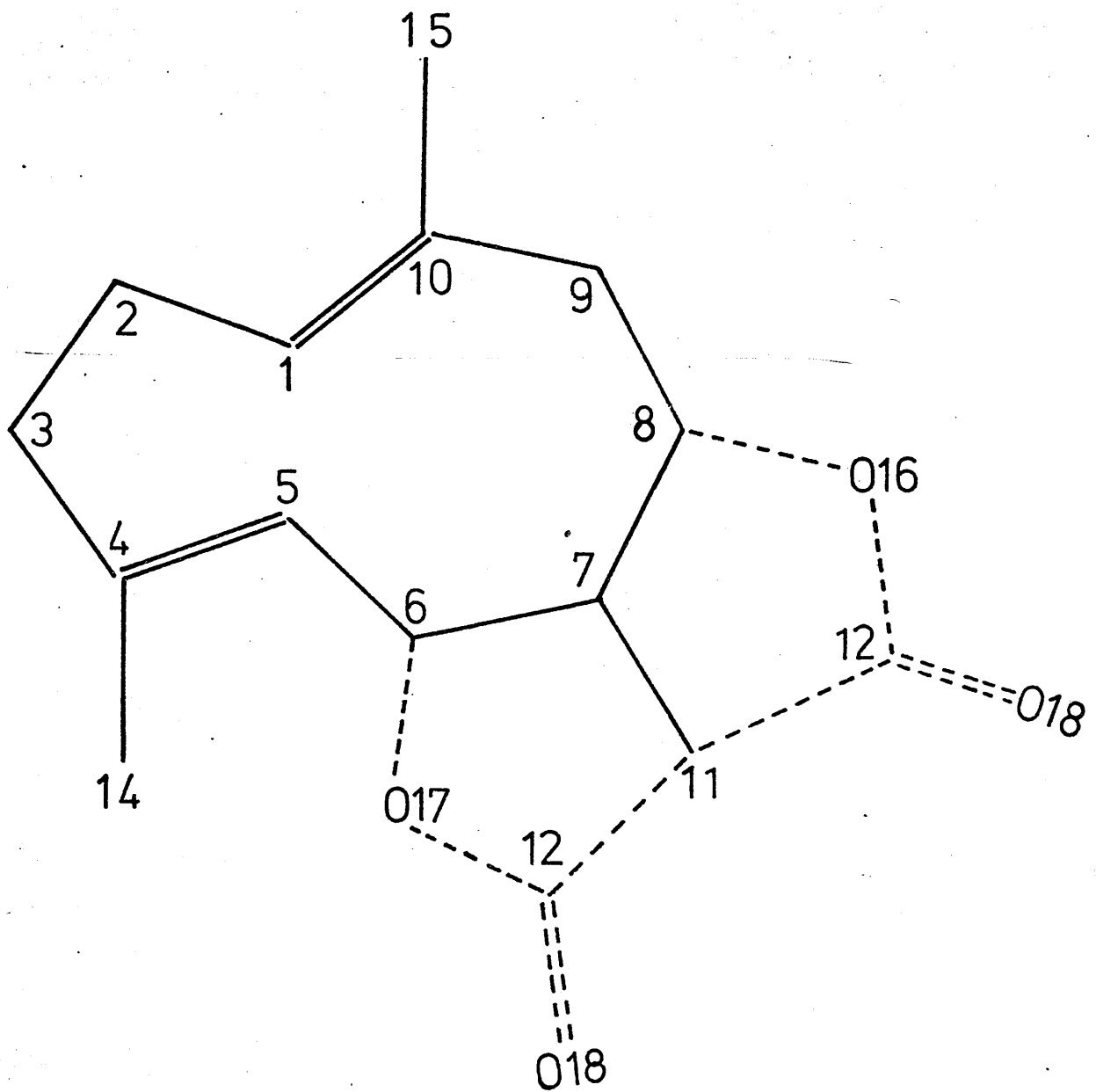
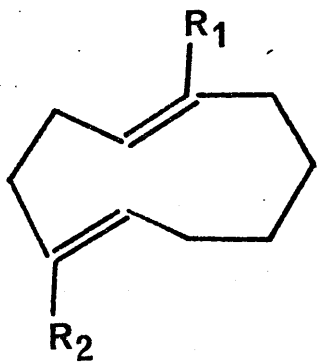
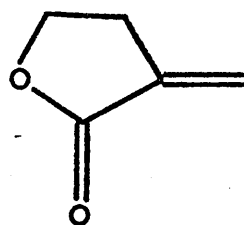


Fig. 6.1 Basic Structure of Germacrane Sesquiterpene γ -Lactones

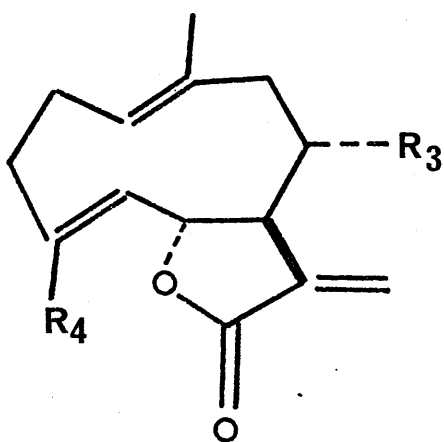


I $R_1 = R_2 = H$

X $R_1 = R_2 = Me$



XIV



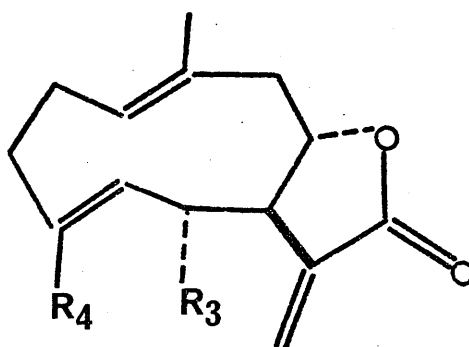
II $R_3 = OH, R_4 = CH_2OH$

III $R_3 = \begin{array}{c} \text{OCOCCH(OH)CH}_2\text{OH,} \\ \parallel \\ \text{CH}_2 \end{array}$

$R_4 = CH_2OH$

IX $R_3 = H, R_4 = Me$

XII $R_3 = OH, R_4 = Me$



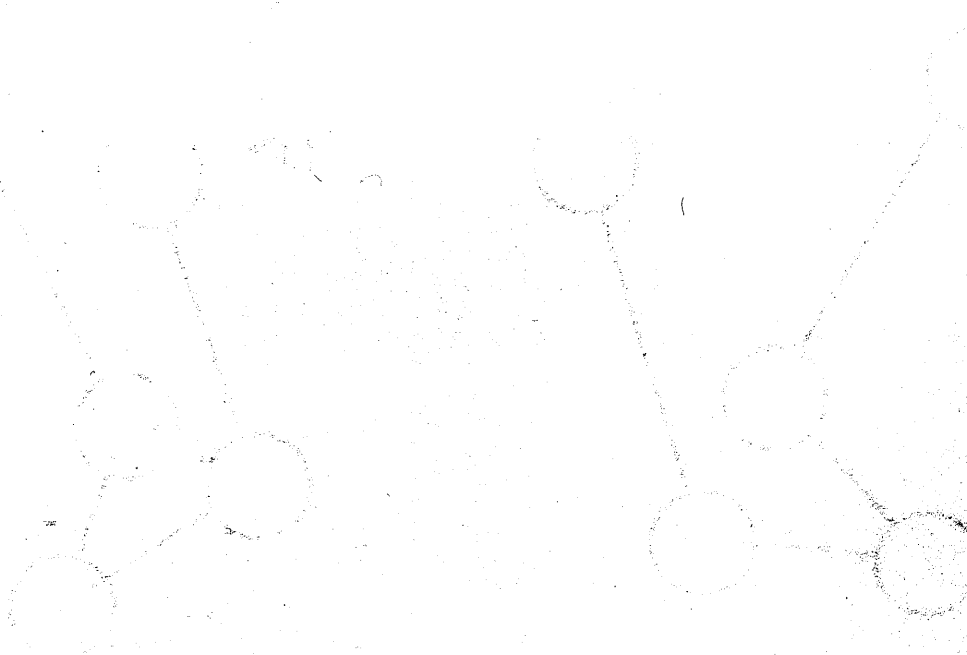
IV $R_3 = OH, R_4 = CH_2OH$

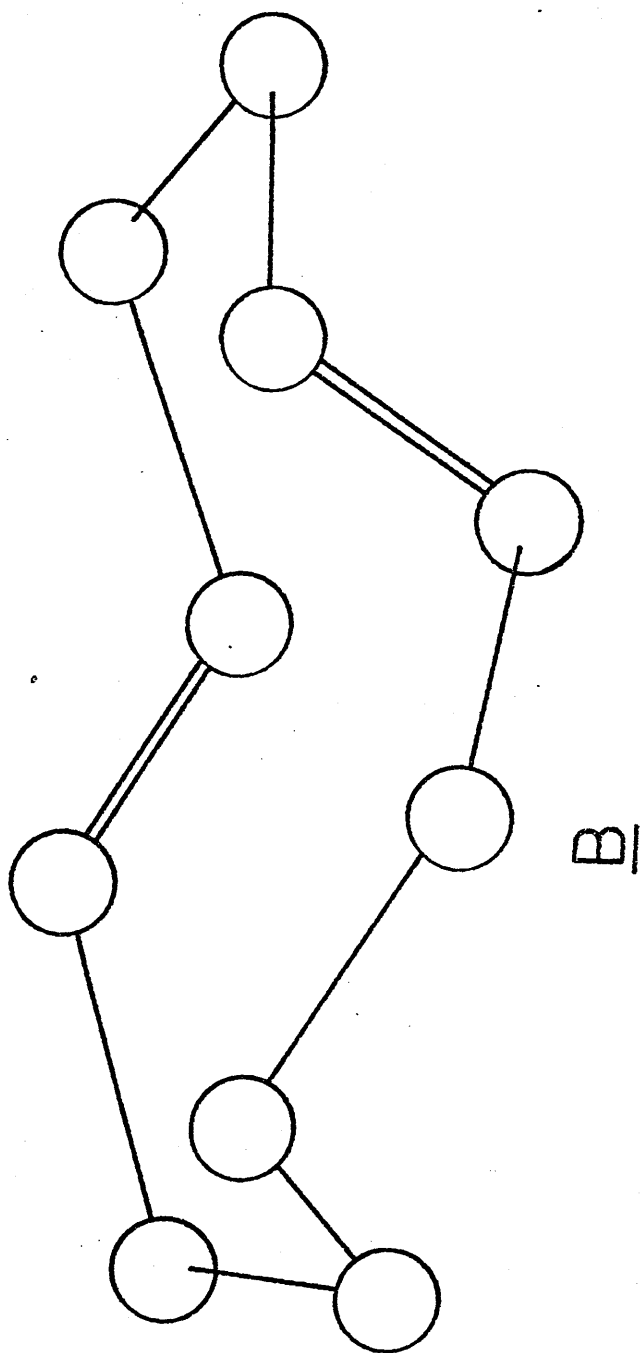
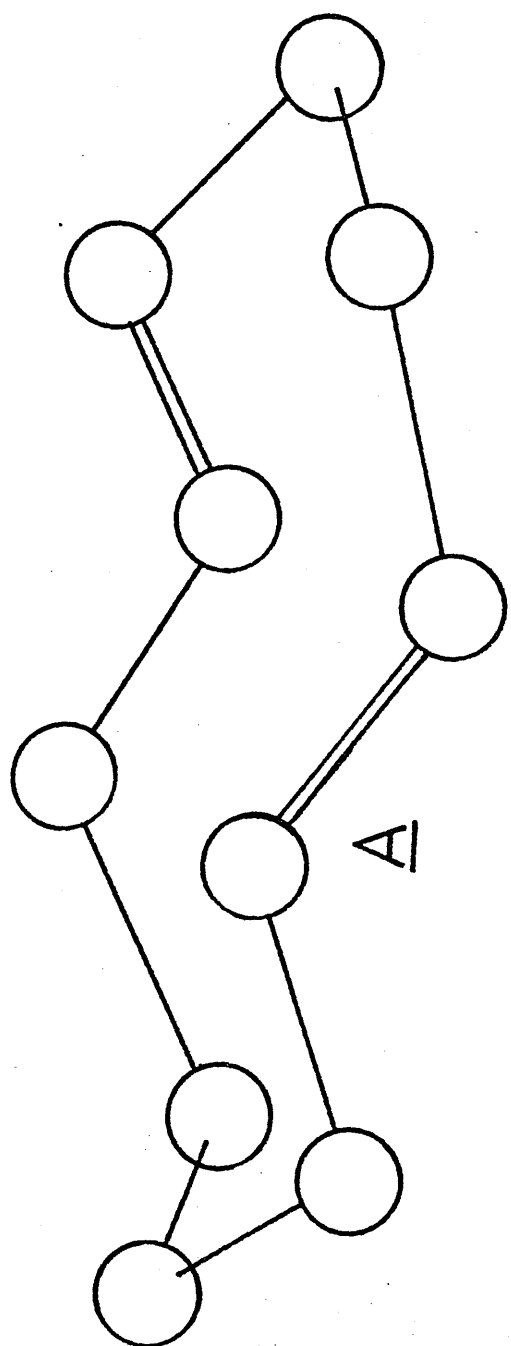
XI $R_3 = H, R_4 = Me$

XIII $R_3 = OH, R_4 = Me$

Fig. 6.2

The A and B type conformations of cyclodeca-1,5-diene





whether it is a feature of the isolated cyclodecadiene.

The location and conformation of the lactone ring poses further questions. Several workers, for instance, noticed that the lactone function was capable of taking part in a rearrangement process in which it could change its fusion from C(6), C(7)— to C(7), C(8)— (Fig. 6.1) (e.g. 11,12). Yoshioka et al (13) on further investigation of trans - fused germacranolides which contain C(6)- and C(8)- lactonisable α -orientated oxygen substitution found that they preferentially relactonise at C(8) after treatment with strong alkali and subsequent acidification. Thus salonitenolide (II) and cnicin (III), for example, form artemisiifolin (IV). The complexity of interatomic forces involved in determining the relative stabilities of the C(6), C(7)— and C(7), C(8)— modes of fusion can only be unambiguously rationalised by a quantitative method such as reliable strain energy calculations.

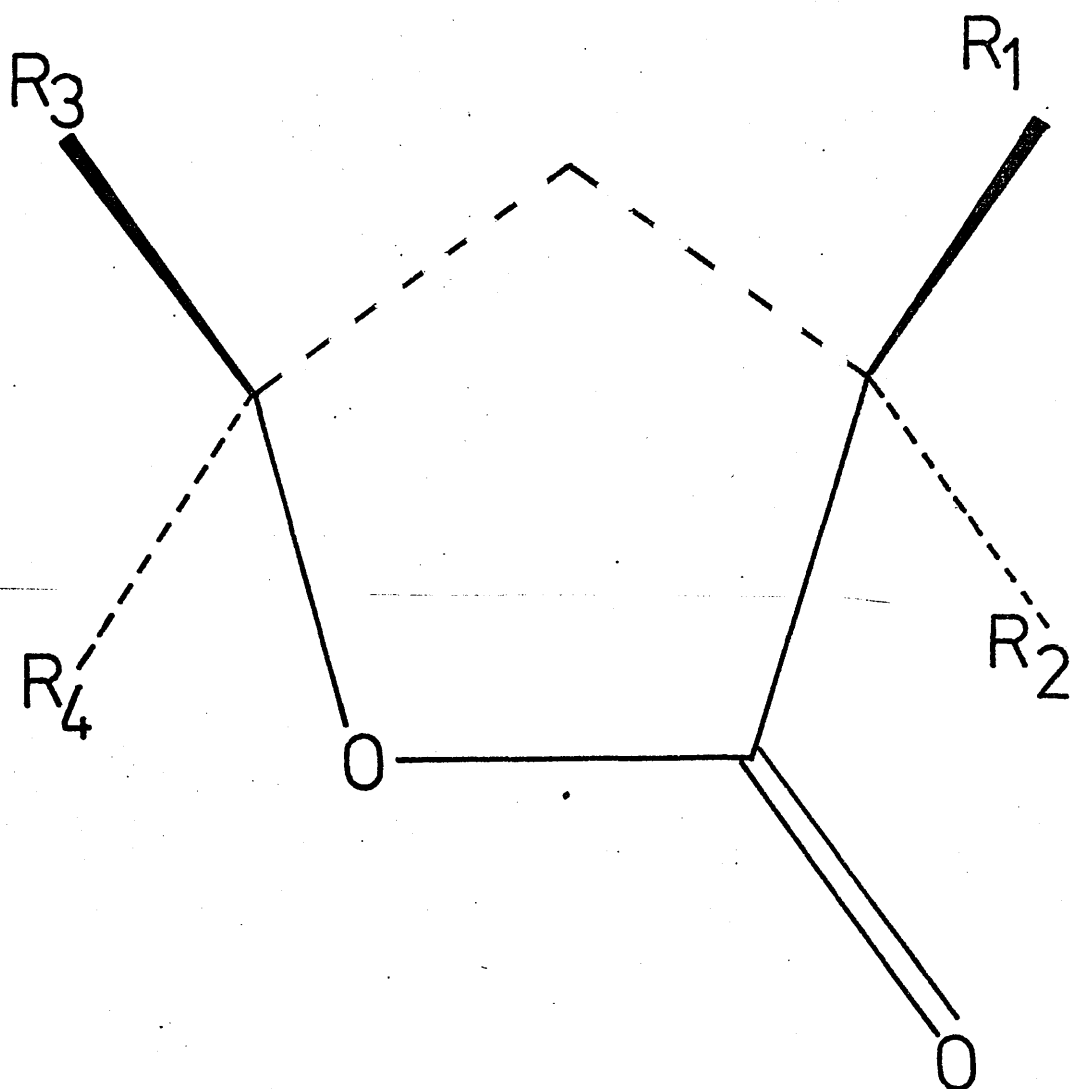
The lactone fusion in germacranolides has been correlated with the Cotton effect at about 250 nm by Beecham (14) who proposed that the sign of this spectrum is related to the chirality of the **C=C-C=O** chromophore. McPhail and Sim (6), however, have noted the apparent insensitivity of the Cotton effect to changes in the **C=C-C=O** torsion angle (Ψ) and have suggested that the torsion angle (Φ) in the vicinity of ring fusion may be more significant. It has also been shown that the value of Ψ can sometimes be significant in determining the relative stabilities of isomeric sesquiterpenes such as α - and β -santonin (16).

The correlation (if any) between $\underline{\Psi}$ and $\underline{\Phi}$ is not immediately obvious and again this is the type of problem which MM should be able to answer. The MM technique has also been used to investigate the relationship between $\underline{\Psi}$ and steric strain in germacranolides.

6.2 Method of calculation

One of the difficulties in using MM to study the problems discussed in section 6.1 is parameterisation of the lactone terms in the force field (FF). The paucity of experimental data for lactones is evident from the inclusion of only one lactone $\Delta H_f^O(g)$ in Cox and Pilcher (18) and a complete absence of electron diffraction results for lactones in Specialist Periodical Reports (19). The force field (FFIII, appendix C) had therefore to be constructed on an ad hoc basis. Fortunately, most of the required terms were already available in an FF which had proved successful in reproducing and interpreting the observed stabilities of several isomers of the santonin family (16). The alkane, alkene, and carbonyl interactions in this FF were taken from Allinger's FF (20,21) because the Glasgow alkane/alkene FF(22) had not been fully parameterised when the calculations were carried out. Allinger's FF has been tested against experimental data for a number of compounds including some steroids (23,24).

FF III, while not being suitable for predicting the absolute strain energies in unrelated compounds should, nevertheless, be adequate for calculating the relative enthalpies of compounds that are structurally similar (25). In order to further test this assertion the FF was applied to the results of some recent equilibration studies on 2,4- disubstituted - γ -butyrolactones (26). These observations yielded the free-energy differences between the cis— and trans isomers of V - VIII and since the entropy differences within each group of geometric isomers are probably very small it is permissible to equate calculated variation in enthalpy with observed free energy. The cis II isomers of V - VIII were not included in the calculations because they are very highly strained as a result of a number of transannular interactions and consequently do not make a significant contribution to the equilibria. Strain energies were computed for the trans I and II and cis I **conformers** of these compounds employing FF III and full matrix minimisation. The results are shown in Table I and because similar energies were obtained for the two trans forms of each compound it was necessary to take a weighted average based on their calculated Boltzmann distribution. The agreement between the calculated and observed results is very good, particularly in view of the relatively small spread in ΔG . The authors of the report on the equilibration studies expressed the view that the cis isomers were more stable because, although the minimum energy form for both cis and trans structures was an envelope in which the endocyclic **C-C(=O)-O-C**



COMPOUND	TRANS I		TRANS II		CIS I		CIS II	
	$R_1=R_4=H$		$R_2=R_3=H$		$R_1=R_3=H$		$R_2=R_4=H$	
	R_2	R_3	R_1	R_4	R_2	R_4	R_1	R_3
V	Me	Me	Me	Me	Me	Me	Me	Me
VI	Et	Me	Et	Me	Et	Me	Et	Me
VII	^t Bu	^t Bu	^t Bu	^t Bu	^t Bu	^t Bu	^t Bu	^t Bu
VIII	ⁿ Bu	Me	ⁿ Bu	Me	ⁿ Bu	Me	ⁿ Bu	Me

TABLE I STABILITIES OF 2,4-DISUBSTITUTED-X-BUTYROLACTONES

COMPOUND	ENERGY *		RATIO **		ENERGY #		DELTA G #	
	TRANS I	TRANS II	TII / TI	TRANS I	CIS I	TRANS I	CALC.†	OBSV.††
V	2.56	2.99	0.49	2.70	2.39		-0.31	-0.14
VI	2.68	2.82	0.79	2.74	2.50		-0.24	0.00
VII	-0.27	-0.05	0.70	-0.18	-1.21		-1.03	-1.20
VIII	2.06	2.23	0.74	2.13	1.86		-0.27	-0.19

K.CAL/MOLE
BOLTZMANN DISTRIBUTION OF ISOMERS AT 298K
WEIGHTED MEAN OF TRANS ENTHALPIES USING RATIO TII / TI
DIFFERENCE (CIS I - MEAN OF TRANS)
FROM REF.26. DELTA G AT 298K

* ** \$ + ††
independent Pbs. together with the measured torsion ...
for conformational ... in Fig. 6.5.

torsion angle was 0° , the cis alkyl (or aryl) substituents were quasi-equatorial, whereas in the trans isomers one of these substituents must be quasi-axial. The calculations fully support this view.

The reliability of FF III is reinforced by its ability to reproduce the geometry of costunolide (IX). The observed (4) and calculated structures of IX are shown in Figs. 6.3 and 6.4 respectively. The root-mean-square (r.m.s.) differences in bond lengths, bond angles and torsion angles between the two determinations are 0.013\AA° , 2.0° and 5° respectively. The corresponding estimated standard deviations in the x-ray study are 0.003\AA° , 0.2° and 0.3° .

The calculations on the butyrolactones and on IX when considered in conjunction with the results reported for the santonins vindicate the use of this FF for investigation of the problems outlined in section 6.1.

6.3 Conformation of the trans, trans - cyclodeca - 1,5 - diene (I)

The frequent occurrence of two of the possible conformations of I in germacranolides was noted in section 6.1. Calculations were carried out on these A and B conformers (Fig. 6.2) using both FF III and the recently parameterised FF I (27). The results from these two independant FFs, together with the measured torsion angles for costunolide (4), are shown in Fig. 6.5.

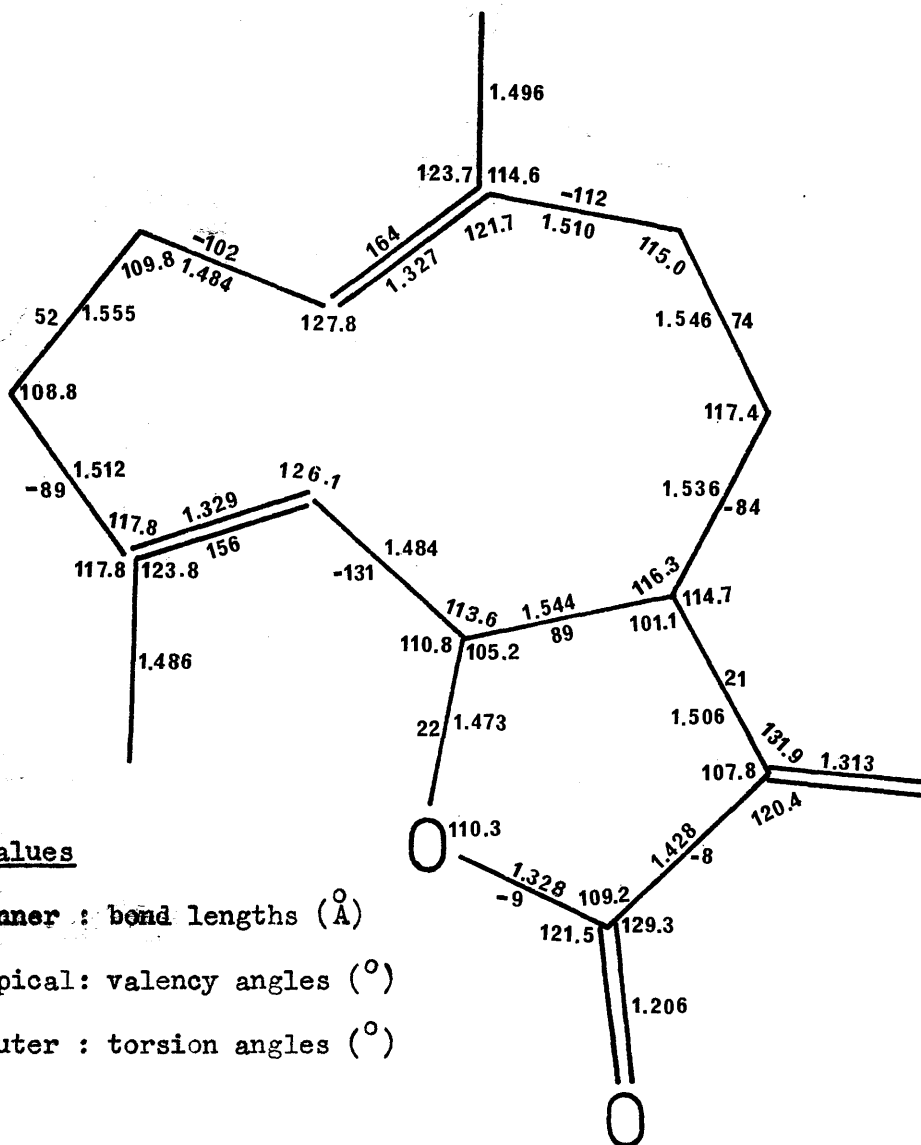


Fig. 6.3 Costunolide: Crystal Structure.

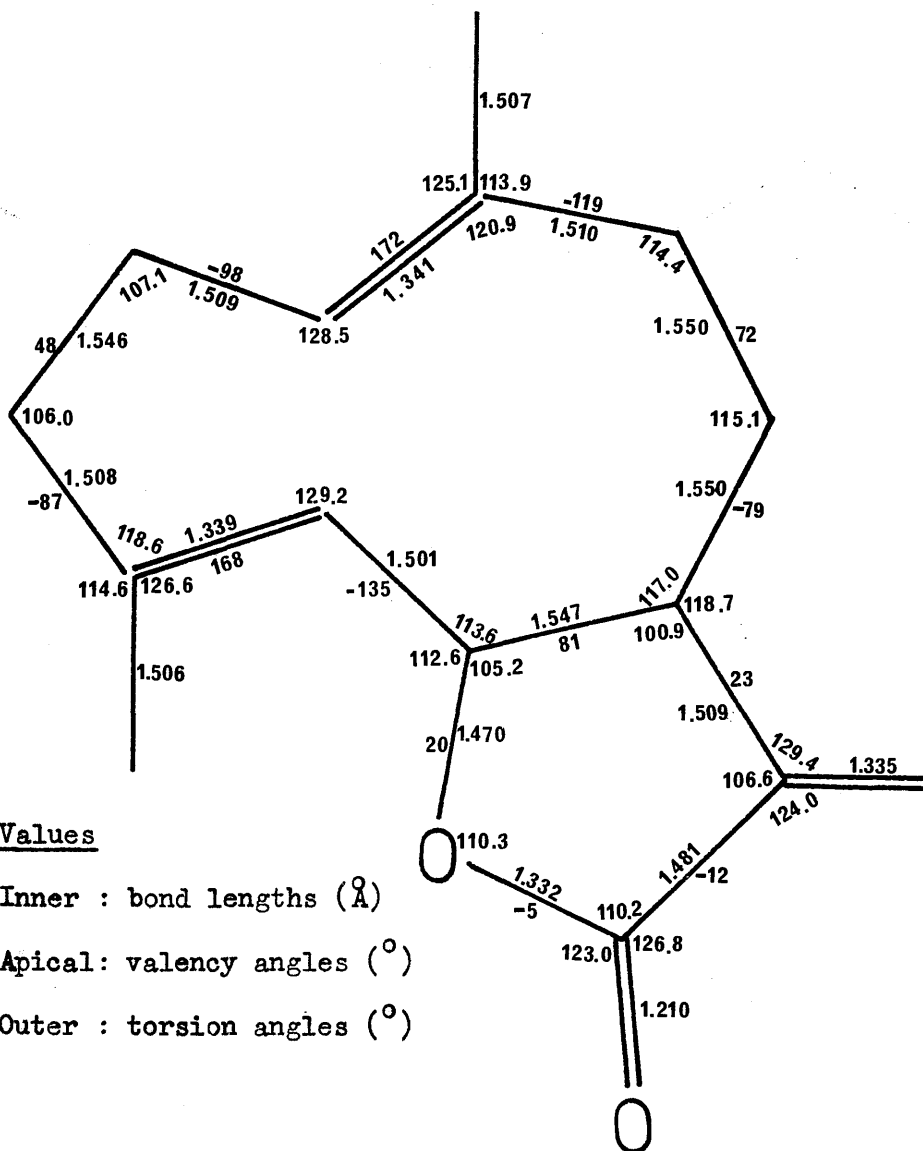
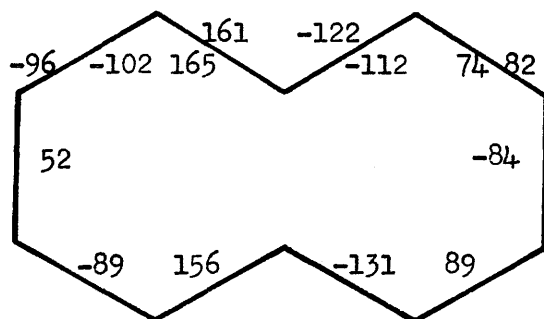
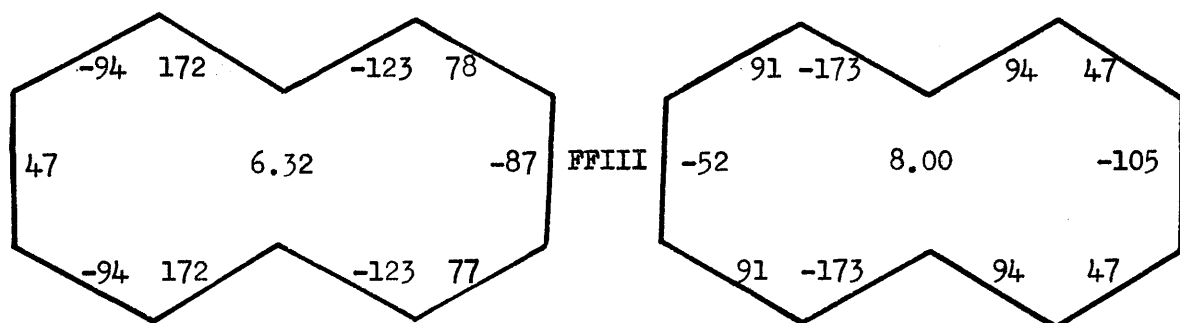


Fig. 6.4 Costunolide: Calculated Structure

A

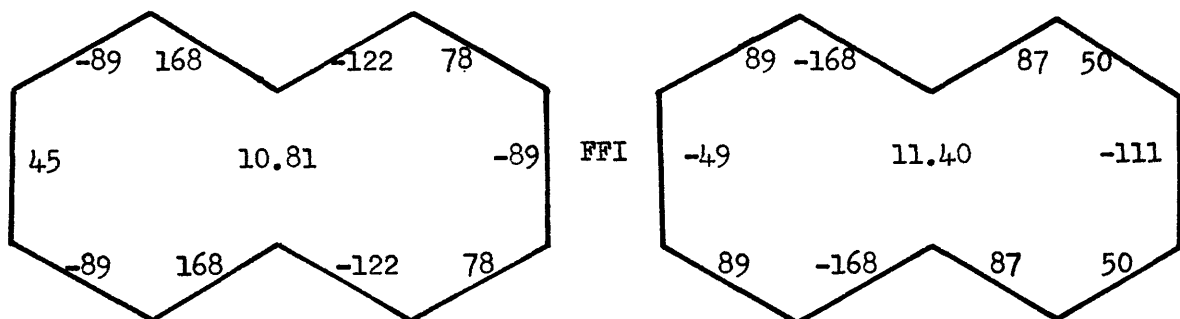
B



Costunolide
Crystal
Structure

A

B



Internal values: Peripheral - torsion angles(degrees)
Central - steric energies(K.cal/mole)
External values: torsion angles averaged to give C₂ symmetry.

Fig. 6.5 Energies and Conformations of A and B type Cyclodeca-1,5-diene

It is encouraging that both FFs give essentially the same stereochemistry and relative stabilities for the A and B conformation of I. This is not entirely unexpected since the main defect of Allingers parameterisation (FF III - alkane/alkene terms), namely overestimation of H ... H non-bonded interactions (28), is not important in these calculations. The conformation adopted by IX in the crystal (Fig. 6.2) is shown to be A and this conformation is about $1.5 \text{ K cal mole}^{-1}$ more stable than B (as typified by shiromodiol). Calculations have also been carried out for a number of other conformations of I using FF I none of which has proved more stable than A or B (27). The unsymmetrical substitution on the cyclodecadiene ring in IX forces a departure from the C_2 symmetry found in the calculation of I. If, however, the measured torsion angles are averaged so as to give a symmetric C_2 conformation of the macrocycle then the calculated and observed A structures lie within the limits of experimental error.

The results of these calculations thus provide a reasonable reference geometry for the two most stable conformations of I and they imply that the unequal deformations about the endocyclic $C-C=C-C$ bonds observed in a number of germacranolides is caused by the substituents to I. In order to test this proposal additional strain energy calculations were performed for IX and X with FF III. The results are shown in Table II and it can be seen that while the torsional deformations have not been fully reproduced by calculation some support is given to the suggestion (4) that the more severe overcrowding

TABLE II CALCULATED AND OBSERVED GEOMETRIES IN THE VICINITY OF THE
DOUBLE BONDS OF A AND B CONFORMATION CYCLODECADIENES.

COMPOUND	TORSION ANGLES *		NON-BONDED SEPARATIONS **			
	A1	A2	D1	D2	D3	D4
I	CONF. A	172	172	2.63	2.87	2.92
X	CONF. A	171	172	2.68	2.78	2.90
IX	CALC.	172	168	2.68	2.76	2.92
IX	X-RAY	164	156	2.83	2.96	3.20
I	CONF. B	173	173	2.72	2.92	2.89
X	CONF. B	172	172	2.64	2.85	3.26
XI	CALC.	172	173	2.64	2.86	3.26
XI	X-RAY +	163	153	2.88	3.03	3.11

CALCULATIONS PERFORMED WITH FFII1

* TORSION ANGLES IN DEGREES
A1 = C(2) - C(1) - C(10) - C(9)
A2 = C(3) - C(4) - C(5) - C(6)

** NON-BONDED SEPARATIONS IN ANGSTROMS
D1 = C(1)...C(4)
D2 = C(1)...C(5)
D3 = C(4)...C(10)
D4 = C(5)...C(10)

THIS IS FOR DIHYDROMIKANOLIDE(7)

about C(4) - C(5) is responsible for the observed consistently greater twisting around this bond in germanolides with skeleton A. Thus the calculated transannular distances for X represent those in the crystal structure much more closely than the corresponding separations in conformation A of I. The substitution of the trans-fused lactone at C(6), C(7) in IX does not significantly change these non-bonded distances. Calculations have also been carried out for XI and the corresponding conformation of X and a similar trend has been observed.

Therefore it seems probable that it is the substitution of the methyl functions at C(4) and C(10) that cause the observed torsional deformations about the Csp^2-Csp^2 bonds in the cyclodecadiene of conformation A. The failure of the FF to reproduce this particular geometrical aspect is disappointing but the exact form of the transannular interaction is so uncertain (4) that it is difficult to devise an FF which will accurately represent the balance of forces in this region. If the difference in the olefinic torsion angles is due to the substitution at C(4) and C(10) then x-ray analysis of derivatives of I (Conformation A) having bulky groups at these locations should also exhibit this deformation. No such analyses have been found in the literature but in 2,7-dibromo-3,8-dimethoxy-trans, trans-cyclodeca-1,6-diene (29) the bromine atoms are displaced outwards from the decadiene ring in a manner not unrelated to that of the methyl functions in IX.

6.4 Preferred fusion and conformation of the lactone ring

The observation of Yoshioka et al (13) regarding the preferred lactonisation of certain sesquiterpene lactones has been investigated by MM. Calculations were carried out for the A and B conformation of compounds IX - XIII and the results (Table III) confirm that the C(7) - C(8) — trans-fused lactone with an oxygen function at C(6) is more stable than the corresponding C(6) - C(7) — fused isomer with an oxygen function at C(8).

The explanation for this behaviour can be found on examination of the steric environment of the lactone ring at each site. Cleavage of the O - CO bond of the δ-lactone of XII or XIII reduces the strain on the cyclodecadiene ring which will relax to a conformation much closer to A or B. It does not appear to be possible for the macrocycle to change from B to the more stable A under these circumstances because the substituents impose severe steric restrictions on such a conformational change.

The major factor influencing the choice of sites now available for relactonisation is the amount of strain arising from the constraints imposed by trans-fusion of the alkene and lactone rings. The minimum energy conformation of an isolated δ-lactone (XIV) was obtained by strain energy calculations (Fig. 6.6). The endocyclic torsion angle about the Csp³ - Csp³ bond of XIV is 20° in its most stable conformation and therefore fusion to the cyclodecadiene ring should be accommodated most easily when the appropriate torsion angle of the ten-membered ring is also about 20°.

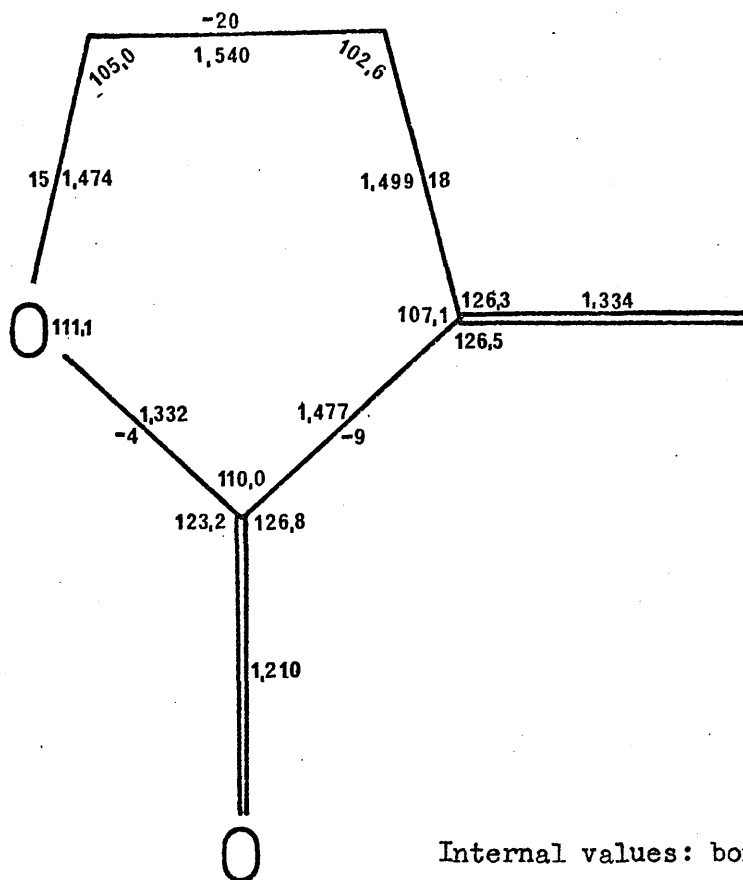
TABLE III STERIC ENERGIES OF SOME GERMACRANOLIDE LACONES

COMPOUND	C(13) METHYLENE		C(13) METHYL +		C(13) METHYL ++	
	A	B	A	B	A	B
ENERGIES						
IX	15.9	20.4	---	---	11.2	---
X	9.1	10.9	9.1	10.9	9.1	10.9
XI	14.6	14.5	9.5	---	---	---
XII	18.7	21.8	12.6	16.1	12.9	19.1
XIII	16.9	16.2	11.5	13.6	11.4	14.6
ENERGY DIFFERENCES						
XII - XIII	1.8	5.6	1.2 *	1.5 *	1.4 **	5.5 **
IX - XI	1.3	5.9	---	---	1.7 **	---

ALL ENERGIES IN K.CAL/MOLE

+ ALPHA METHYL
 ++ BETA METHYL
 * C(13) ALPHA TO BETA
 ** C(13) BETA TO ALPHA

SEE TEXT
 SEE TEXT



Internal values: bond lengths(Å)

Apical values: valence angles(°)

External values: torsion angles(°)

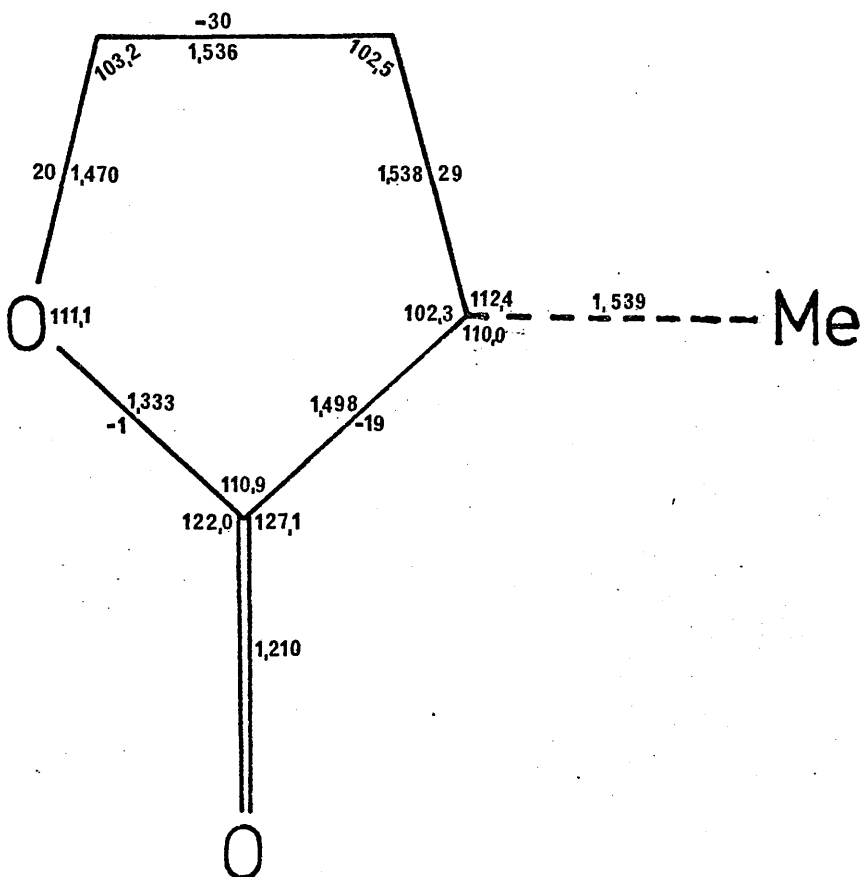


Fig. 6.6 Calculated minimum energy geometries for XIV and one of its C(13)-methyl analogues

Newman projections down (i) the C(6) - C(7) and (ii) the C(7) - C(8) bonds of the calculated structures for the A and B conformations of 1,5-dimethyl-cyclodeca-1,5-diene (X) are illustrated in Fig. 6.7. The locations for subsequent lactone fusion are shown by broken lines. For the B conformations it is quite clear that C(7) - C(8) fusion is to be preferred since the relevant torsion angles are -59° and -3° at (i) and (ii) respectively. Compound XIII therefore represents the minimum energy structure for the B conformation and it is interesting to note that the additional strain is accommodated mainly by the lactone ring deforming so that it is almost planar (Fig. 6.8). However, the planar lactone ring is not as unstable as might be anticipated since the endocyclic valence angles move closer to 108° and this is nearer to their unstrained values ($109 - 120^{\circ}$) than in XIV thus reducing the Baeyer strain. Also the minimum Pitzer strain is at 0° for all the bonds of the lactone except the single $\text{Csp}^3 - \text{Csp}^3$ bond where it is maximised. Although the planar lactone is less favourable than the puckered conformation in isolation it appears that in a gross structure such as the B conformation of XIII it can represent the minimum energy stereochemistry. It has been observed experimentally in the x-ray analyses of elephantol (5) and a derivative of florilenalin (31).

C(7) - C(8) trans-fusion of the α -methylene -Y-lactone is also preferred for the A conformation. The difference between the appropriate torsion angles ((1) -44° , (11) 38°) is not enough to account for the 1.8 K.cal per mole energy differential between the A conformations

Fig. 6.7

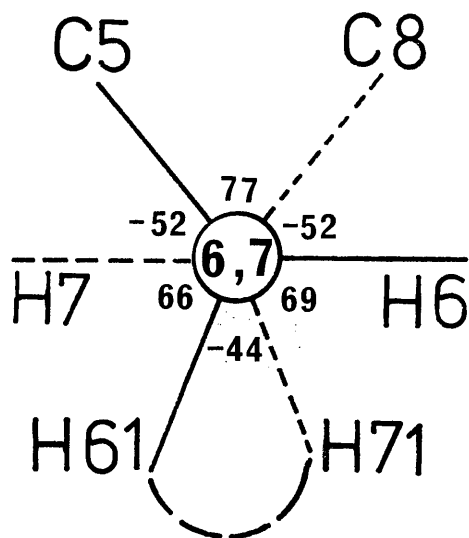
Newman projections down the C(6)-C(7) and C(7)-C(8) bonds
for the calculated structures of the A and B type conformations
of dimethylcyclodeca-1,5-diene.

(a) C(6)-C(7) A type

(b) C(7)-C(8) A type

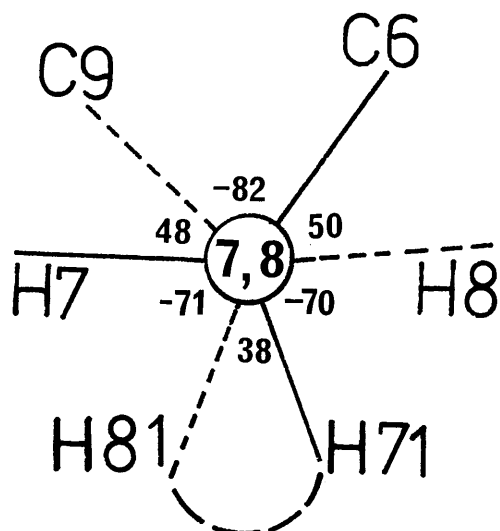
(c) C(6)-C(7) B type

(d) C(7)-C(8) B type

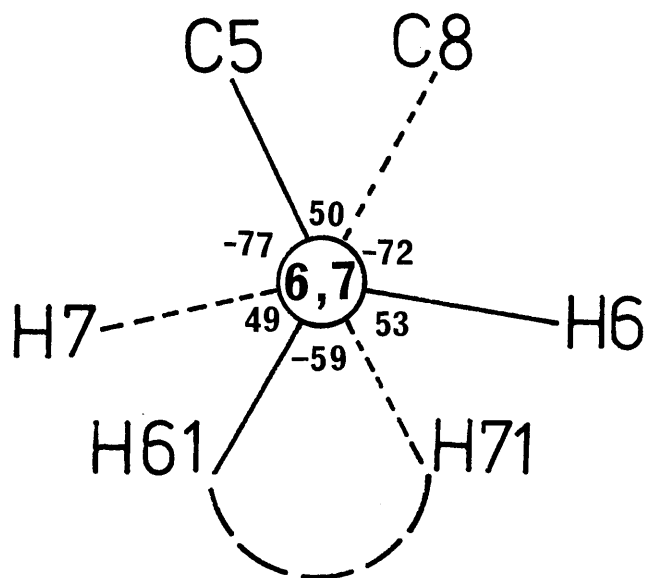


(a)

A

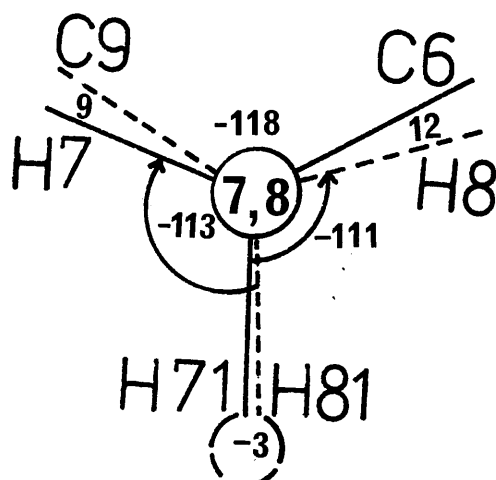


(b)



(c)

B



(d)

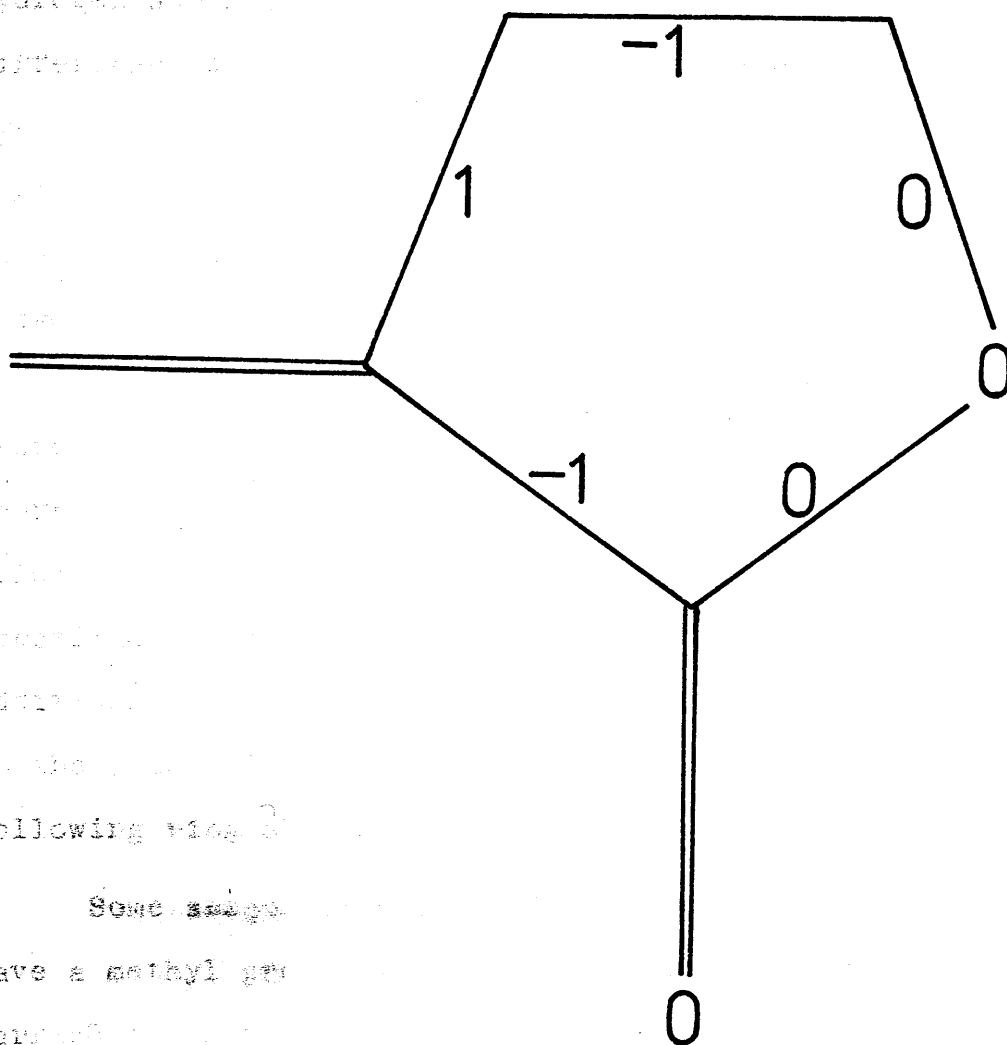


Fig. 6.8 Calculated torsion angles($^{\circ}$) for lactone in XIII (B)

of XII and XIII (Table III). Further investigation has indicated that there is a secondary stabilising effect for the C(7) - C(8) fusion arising from the distinct steric environments of the hydroxyl group at C(6) and C(8). If the hydroxyl functions in XII and XIII are replaced by hydrogen atoms (IX and XI) then the relevant energy difference is 1.4 k cal per mole. Complete removal of the α -hydrogen at C(6) or C(8) in these latter two compounds further reduces this difference to 0.4 k cal per mole. This suggests that the C(6) α -hydroxyl is in a less crowded environment than the α -hydroxyl at C(8). The main reason for this is the interaction between the C(8) hydroxyl and the hydrogens attached to C(9) (Table IV). There is only one hydrogen at the sp^2 C(5) so that this effect is considerably reduced for the C(6) hydroxyl substituent and consequently relactonisation at C(8) is additionally stabilised. There is a much smaller difference in the alternative environments of the ester oxygen atom following ring closure.

Some sesquiterpene lactones (e.g. dihydromikanolide) have a methyl group at C(11) and calculations have been carried out for the corresponding α -methyl isolated lactone (Fig. 6.6) and for the A and B conformations of IX - XIII with α - and β -methyl substitution at C(11) to confirm the generality of the preferred C(8) relactonisation. When the methylene group at C(11) is replaced by a methyl function, rotation about the C(11) - C(13) bond can relieve strain and thus reduce the H(C13) ... O(hydroxyl) interaction (Table IV). Also C(11) is now sp^3 changing the

TABLE IV CONFORMATION A: SOME NON-BONDED INTERACTIONS INVOLVING THE
HYDROXYL FUNCTION

FUSION			
C (6) - C (7)		C (7) - C (8)	
ATOMS	SEPARATION *	ATOMS	SEPARATION *
C (9) ... H (O16)	2.53	C (5) ... H (O17)	2.47
O (16) ... H (C7)	2.69	O (17) ... H (C7)	2.74
O (16) ... H (C9)	2.63	O (17) ... H (C5)	2.75
O (16) ... H* (C9)	2.63	---	---
O (16) ... H (C13)	2.30	O (17) ... H (C13)	2.32
H (C8) ... H (O16)	2.20	H (C6) ... H (O17)	2.23
H* (C9) ... H (O16)	2.34	H (C5) ... H (O17)	2.75

* IN ANGSTROMS

O (16) - H (O16) IS THE HYDROXYL GROUP AT C (8) IN C (6) - C (7) FUSED LACTONES
O (17) - H (O17) IS THE HYDROXYL GROUP AT C (6) IN C (7) - C (8) FUSED LACTONES

INTERACTIONS READING ACROSS THE PAGE ARE EQUIVALENT.
THUS C (9) ... H (O16) IS EQUIVALENT TO C (5) ... H (O17) FOR C (7) - C (8) FUSION

minimum of several torsional interactions from 0° to 60° . The net result is that the isolated α -methyl lactone is more stable (than the methylene) by 3.0 K.cal per mole. The calculated energies for the gross structures (Table III) show that C(7) - C(8) fusion is once again more favourable, although the relevant energy differences are generally smaller. Comparison of isomers is a little more complicated because, for instance, a C(6) - C(7) fused compound with C(13) α -methyl would, on relactonisation to C(7) - C(8) fusion have the C(13) methyl β (Fig. 6.9). The energy differences quoted in Table III allow for this additional constraint.

The experimental work of Yoshioka et al is therefore supported by these strain energy calculations which extend the range of compounds to which the relactonisation rule can be applied.

The preceding discussion on lactone fusions in germacranolides does not consider the possibility of cis-fusion. If it is assumed as in all known germacranolides, that C(7) - C(11) is always β then H(6) and H(8) (Fig. 6.7) represent the alternative locations at which lactonisation could occur. For conformation A of X the appropriate torsion angles of 69° and -70° , for C(6) and C(8) fusion respectively, are very much further away from the minimum conformation of the isolated lactone than the torsion angles which would be involved in trans-fusion. This is reflected in the higher strain energy of 20.6 K.cal per mole for the cis-lactone isomer of XII compared

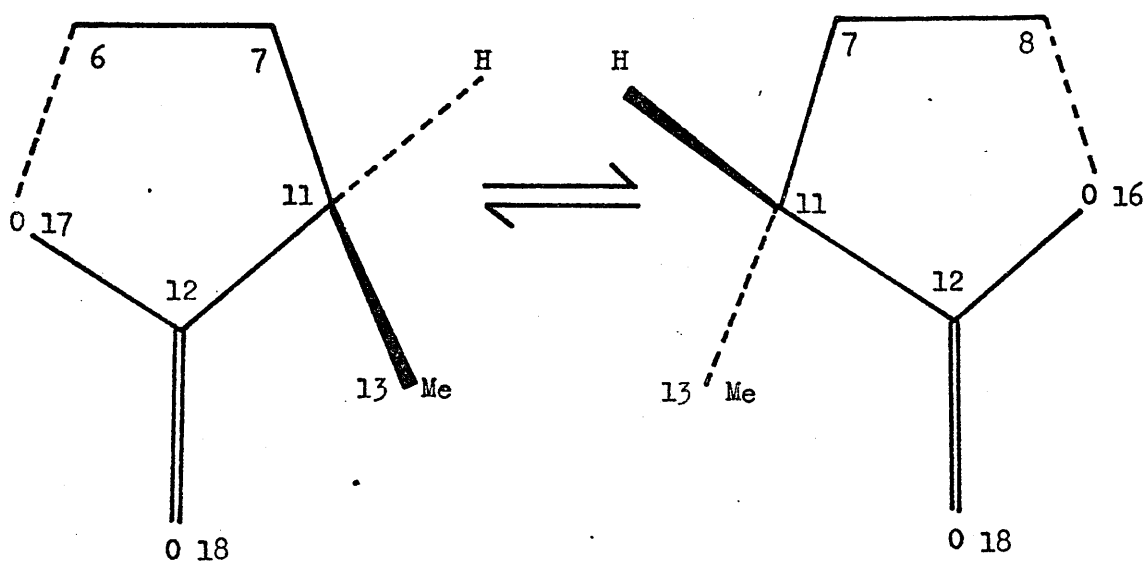
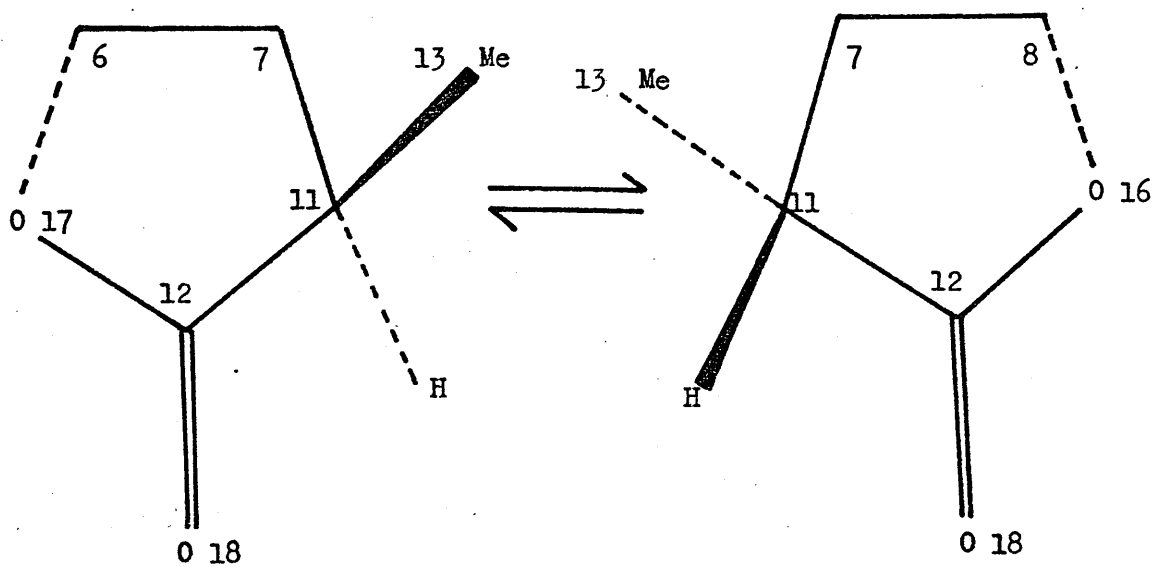


Fig. 6.9 Relationships between stereochemistry at C(11) and lactone fusion

with the corresponding value of 18.7 Kcal per mole for the trans-fused lactone.

In the B conformation of X cis-fusion at C(8) is extremely unfavourable (Fig. 6.7) and the much smaller value of the relevant torsion angle at C(6) (59° as opposed to -111°) makes this latter site the most acceptable.

The proposals made in this section regarding the energetically favoured sites for lactone fusion are based on the assumption that the lactone ring retains a geometry close to that of the calculated energy minimum. This supposition was tested by applying the algorithm of Cremer and Pople (32) to a number of calculated and observed structures. The results in terms of their parameter \varnothing are listed in Table V, and show that there is a remarkable consistency in the conformation of the lactone ring in a range of very different germacranolides. This conformation appears, for the examples quoted, to be independent of:

- (i) The mode of fusion. Unfortunately the very unfavourable 7,8 cis fusion is not represented but the 6,7 cis- and the trans- fused lactones show little variation in conformation.
- (ii) The group substituted at C(11) even when there is a very bulky group at C(8) e.g. in Eupatoriopicrin (34).
- (iii) The conformation of the macrocycle of the germacranolide.

The values listed in Table V also indicate the location of the conformation on the pseudorotational

TABLE V CONFORMATION OF THE LACTONE RING IN SOME GERMACRANOLIDES

COMPOUND	TYPE C (13)	ALPHA	PHI	A\$	GAM	PHI	B\$	GAM
	*	***	##	PSI		##	PSI	
<u>CALCULATIONS</u>								
ISOLATED LACTONE \$	---	---	10	-9	-20	---	---	---
ISOLATED LACTONE \$	---	---	1	---	-30	---	---	---
IX	6,7T	H	10	-11	-26	11	-12	-29
	6,7T	H	1	---	---	---	---	---
XII	6,7T	OH	8	-15	-29	13	-9	-26
	6,7T	OH	2	---	-30	-1	---	---
	6,7T	OH	-3	---	-33	1	---	---
XI	7,8T	H	10#	10	23	PL	0	0
	7,8T	H	1#	---	---	---	---	---
XIII	7,8T	OH	9#	13	27	PL	-1	-1
	7,8T	OH	-1#	---	32	-14#	---	12
	7,8T	OH	0#	---	32	-9	---	-16
	6,7C	OH	9#	16	34	---	---	---
<u>CRYSTAL STRUCTURES</u>								
COSTUNOLIDE (4)	6,7T	H	17	-10	-25	---	---	---
ALATOLIDE (6)	6,7T	OXY	23	0	-19	---	---	---
EUPATOLIDE (34)	6,7T	OXY	15	-12	-27	---	---	---
EUPATORIOPICRIN (35)	6,7T	OXY	5	-9	-24	---	---	---
ONOPODIOPICRINE (35)	6,7T	OXY	11	1	-16	---	---	---
DIHYDROMIKANOLIDE (7)	7,8T	OXY	---	---	---	1#	---	31
ELEPHANTOL (5)	7,8T	OXY	---	---	---	PL	11	33

SYMBOLS EXPLAINED ON FOLLOWING PAGE

EXPLANATION OF SYMBOLS IN TABLE V

OF FUSION OF LACTONE TO CYCLODECADIENE
 E.G. 6,7T MEANS C(6) - C(7) TRANS-FUSED
 E.G. 7,8C MEANS C(7) - C(8) CIS-FUSED
 SUBSTITUTION AT C(11)
 M1 C(13) IS =CH2
 M2 C(13) IS ALPHA METHYL
 M3 C(13) IS BETA METHYL
 SUBSTITUTION AT THE ALPHA POSITION OF C(6)/C(8) FOR 6,7/7,8 FUSION

PHI (DEGREES) CALCULATED WITH ALGORITHM OF CREMER AND POPL (32)
 BASE VALUE (0) IS THE ALPHA-C(7) ENVELOPE
 INTERVAL BETWEEN ENVELOPES/HALF CHAIRS IS 18 DEGREES
 GAM (DEGREES) IS THE ENDOCYCLIC C-C-C-O TORSION ANGLE OF
 THE LACTONE RING
 PSI (DEGREES) IS THE EXOCYCLIC C=C-C=O TORSION ANGLE OF
 THE LACTONE RING
 RESULTS FOR ISOLATED LACTONES GIVEN UNDER A CONFORMATION FOR
 CONVENIENCE

PHI IS EQUAL TO OR GREATER THAN 180 DEGREES THEREFORE ROTATION IS
 FROM BETA-C(7) ENVELOPE
 SIGN OF PHI GIVES ROTATION FROM C(7) ENVELOPE
 + IS ANTICLOCKWISE 6,7 FUSION
 - IS CLOCKWISE 6,7 FUSION
 + IS CLOCKWISE 7,8 FUSION
 - IS ANTICLOCKWISE 7,8 FUSION

PLANAR LACTONE RING
 NON-HYDROXYL OXYGEN FUNCTION E.G. ESTER

show, in this table and the reported correlation (33) between these two angles is observed to all the examples

circuit. Thus in an isolated 5-membered cycle it is possible to have the puckering (e.g. for an envelope) α or β -orientated so that a calculated geometry for an isolated cyclopentane is not unique because there is an isoenergetic isomer in which all the torsion angles have their signs reversed. In the isolated γ -lactone with methylene substitution at C(11) (Fig. 6.6) the mirror image is also isoenergetic and the five-membered ring can adopt α or β puckering with equal ease. When the C(11) substitution is methyl (M2 or M3 in Table V) there will be a small difference in stability between the mirror-image related conformations. The results in Table V show that the lactone conformation in germacranolides generally approximates to a C(7) envelope (in this context this includes the half-chairs in which C(7) is one of the out of plane atoms). The major exception to this being the occurrence of planar stereochemistry when the conformation of the macrocycle makes this more stable.

Further, 6,7 trans-fused germacranolides should adopt a C(7) α -puckered envelope conformation of the five-membered ring independent of the conformation (A or B) of the macrocycle while the 7,8 trans-fused isomer (A conformation) should be β . The A conformation crystal structures listed in Table V all obey this rule.

The values of the lactone ring torsion angles ψ (C=C—C=O) and ϕ (endocyclic C—C—C—O) are also shown in this table and the reported correlation (33) between these two angles is observed in all the compounds investigated. ψ and ϕ are also related to the Cremer and Pople angle θ and the sign of ϕ (+ or -) can be used as a

predictor of the lactone puckering ($\underline{\beta}$ or $\underline{\alpha}$ respectively). Examination of Dreiding models indicates that, for lactone conformations close to a C(7) envelope, this latter correlation is exactly what would be expected.

The reasons for the relationship between the lactone puckering and the mode of fusion (6,7 trans or 7,8 trans) with the dimethyl cyclodecadiene can be elucidated from Fig. 6.7. The minimum energy conformation of the isolated lactone XIV (Fig. 6.6) has $\Phi = -20$ and the existence of an isoenergetic isomer with $\Phi = +20$ has been noted above. The stereochemistry about the C(6) - C(7) bonds of both A and B conformations of X is very similar and in particular the H(61)—C(6)—C(7)—H(71) torsion angles are negative (-44° and -59° respectively) and hence 6,7 trans lactonisation is facilitated with the $\Phi = -20$ enantiomer resulting in α puckering. The relevant torsion angle for 7,8 trans-lactonisation for conformation A is 38° resulting in β puckering. The corresponding torsion angle in the B conformation is -3° and the lactone ring is able to adopt α , β , or planar stereochemistry. (Table V). On this basis prediction can also be made for the cis-fused lactones, which have so far not been observed in nature, by considering the H(6)—C(6)—C(7)—H(71) torsion angles of X. For instance a 6,7 cis fused germacranolide with an A conformation of X should be β puckered. The calculated minimum energy conformation of such a structure has a β puckered lactone ring (Table V). It should also be possible to extend this approach to account for lactone conformations in other sesquiterpenes once the ground state of the main carbon skeleton has been

TABLE VI SUMMARY OF RESULTS IN SEC. 6.4

PREFERRED FUSION OF LACTONE	CONFORMATION A 7,8 TRANS	CONFORMATION B 7,8 TRANS
LACTONE STEREOCHEMISTRY		
ENDOCYCLIC C-C-C-O TORSION ANGLE	+	BETA PUCKER
ENDOCYCLIC C-C-C-O TORSION ANGLE	-	ALPHA PUCKER
CORRELATION BETWEEN LACTONE STEREOCHEMISTRY, MODE OF FUSION, AND CYCLODECADIENE CONFORMATION		
6,7 TRANS	CONFORMATION A ALPHA	CONFORMATION B ALPHA
6,7 CIS	BETA	BETA
7,8 TRANS	BETA	*
7,8 CIS	ALPHA	ALPHA

THESE RESULTS ASSUME THAT THE LACTONE CONFORMATION IS CLOSE TO THE MINIMUM ENERGY STEREOCHEMISTRY OF THE ISOLATED LACTONE AND THAT THE C(7) - C(11) BOND IS ALWAYS BETA

* CAN BE ALPHA, BETA, OR PLANAR. SEE TEXT.

obtained. The substitution at C(11) (methyl or methylene) appears to make very little difference to the puckering of the lactone ring except when the relevant torsion angle on the larger carbocycle is close to 0° (as for 7,8 fusion of B conformation germacranolides.)

The conclusions made in this section are summarised in Table VI.

6.5 The $C=C-C=O$ chromophore

The relationship between the stereochemistry of lactone ring fusion and the sign of the Cotton effect at 250 - 260 nm, and also its apparent correlation with the endocyclic $C-C-C-O$ (ϕ) and exocyclic $C=C-C=O$ (ψ) torsion angles has been noted previously (e.g. 4, 6, 15, 36). Cox and Sim (6, 33) have questioned the relevance of the Beecham correlation (15) between the sign of the Cotton effect and the chirality of the $C=C-C=O$ chromophore as a result of the observed value ($\psi = 0$) obtained for alatolide. The recent determination of the structure of onopordopicrine found a ψ of 1° with an estimated error of 0.5° (34) giving rise to further doubts since Beecham predicted that such compounds should have left handed chirality of the $C=C-C=O$ group (ψ negative). Further, the results in the previous section support the view of Cox and Sim, since a direct correspondence has been shown between ϕ and the mode of lactone fusion for A and B type germacranolides. Thus there is a relationship between the sign of ϕ (as assigned by Cox and Sim (6), McPhail and Sim (33)), the

correlation of Stöcklin et al (15) between the sign of the Cotton effect and the stereochemistry and location of lactone fusion, and the prediction of Section 6.4 (Table VII).

The value of $\underline{\Psi}$ appears to be less important than $\underline{\Phi}$ and the wide range in the magnitude of $\underline{\Psi}$ suggests a degree of flexibility about the C(11) - C(12) bond.

In the isolated molecule XIV the minimum stereochemistry is a C(7) envelope (Fig. 6.6) and the isoenergetic $\underline{\alpha}$ - and $\underline{\beta}$ -envelopes are observed (from Dreiding models) to have $\underline{\Phi}$ and $\underline{\Psi}$ paired in sign. The calculation of the $\underline{\alpha}$ -envelope, for instance, gives values of -20° and -9° respectively. Thus to a first approximation the chiralities are interdependent. The minimum energy XIV still has some bad van der Waals repulsions (C(13)...H(7¹), 2.82\AA° ; H(7)...H(6¹), 2.43\AA° ; and C(13)...O(18), 3.00\AA°) which may become more severe in a sesquiterpene. The angle $\underline{\Psi}$ has the minimum torsional interaction at 0° and the value of -9° in the isolated lactone is caused by an attempt to minimize the two bad C(13) non-bonded interactions (H(7¹) and O(18) are directed towards opposite faces of the lactone). Clearly the variation of steric energy with $\underline{\Psi}$ in a sesquiterpene is dependant on the degree to which these (and other) interactions can be minimized in a specific environment and this should be reflected in the range of observed $\underline{\Psi}$.

The method of Wiberg and Boyd (37) has been used to pseudorotate $\underline{\Psi}$ at 3° intervals for the A and B conformations of XII and XIII and the results are plotted in Fig. 6.10. With the exception of the minima, the calculations were

TABLE VII RELATIONSHIPS BETWEEN THE LACTONE STEREOCHEMISTRY AND THE COILION EFFECT AT 260N.M.

LACTONE FUSION	SIGN OF		SIGN OF GAMMA		G \$
	CE	*	CS & MS	+	
6,7 TRANS	-		-		-
7,8 TRANS	+		+		+
6,7 CIS	+				+
7,8 CIS	-				-

GAMMA IS THE ENDOCYCLIC C-C-C-O TORSION ANGLE

* CORRELATION OF STOCKLIN ET AL(14)

+ AS ASSIGNED BY COX AND SIM(6), MCPHAIL AND SIM(33)

\$ AS REPORTED IN THIS THESIS. APPLICABLE ONLY TO THE A AND B CONFORMATIONS OF GERMACRANOLIDES

Original: Energy

Ab initio: Endocyclic C-C-C-O torsion angle (psi)

Conformation

- 324 -

Conformation

A

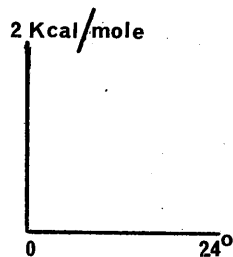
Lactone Fusion

B

6,7 trans

7,8 trans

6,7 cis



Ordinate: Energy

Abcissa : Exocyclic C=C-C=O torsion angle (psi)

Fig. 6.10 Variation in energy with psi for various germacranolides

performed to block diagonal convergence only (17). In germacranolides with the A conformation $|\underline{\psi}|$ varies by about 15° (Table V) and the calculated increase in steric energy at $\pm 15^\circ$ from the minimum in each of the calculated structures is given in Table VIII. This table also includes the results for a 6,7 cis fused A type germacranolide.

There is a correspondence between the flexibility in $\underline{\psi}$ and the ease or otherwise of lactonisation of the appropriate dimethyl cyclodecadiene (Section 6.4). The strain present in a 6,7 trans fused B conformation— and — 6,7 cis fused A conformation— is clearly reflected in the additional strain induced when $\underline{\psi}$ is removed from its value in the minimum energy structure (as shown by the steeper-sided potential wells for these structures in Fig. 6.10). In the other compounds a certain amount of flexibility is apparent and it is also evident that for the A conformations, in particular, variation of $\underline{\psi}$ in one direction (positive or negative) is more favourable than in the other. Thus it would be expected that for 6,7 trans fused A type germacranolides most compounds would have $\underline{\psi}$ within the range 0 to -15° ($\underline{\psi}^+ = 0.4$, $\underline{\psi}_{\min} = -15$).

The observed $\underline{\psi}$ in the corresponding structures in Table V ranges from -10 to $+1$. The energy difference between the minimum and one of the extrema is about $1 \text{ K.cal mole}^{-1}$ for most of the structures in Table VIII.

TABLE VIII VARIATION OF STERIC ENERGY WITH PSI

COMPOUND	PSI (MIN)	RELATIVE ENERGIES \$			K.CAL/MOLE	THIS IS THE CIS FUSED ISOMER OF VIII	INTERPOLATED VALUE. STEEP GRADIENT TO CURVE AND CALCULATION DID NOT EXTEND TO 15 DEGREES FROM MINIMUM	IS THE EXOCYCLIC C=C-C=O TORSION ANGLE (DEGREES)	IS THE VALUE OF PSI IN THE MINIMUM ENERGY STRUCTURE	DIFFERENCE BETWEEN STRAIN ENERGY AT PSI (MIN) AND AT PSI (MIN + 15)	DIFFERENCE BETWEEN STRAIN ENERGY AT PSI (MIN) AND AT PSI (MIN - 15)
		PSI +	PSI -	PSI +							
VIII	-15	0.4	1.5	>5.0 *	2.0						
IX	13	1.5	0.8	1.1	1.3						
VIII (CIS)	16	2.0	3.2 *								

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6.6 References

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APPENDIX A

FFI GLASGOW ALKANE/ALKENE FORCE FIELD

ATOM KEY

1 = H, 2 = C(SP)**2, 3 = C(SP)**3,

FORCE CONSTANTS ARE IN KCAL/MOLE/ANGSTROM**2 OR IN KCAL/MOLE/DEGREE**2
ENERGIES ARE IN KCAL/MOLE

GLASGOW ALKANE/ALKENE FORCE FIELD PAGE NUMBER 2

BOND STRETCHING

EXPRESSION: EB=STRETCHK1*(LENGTH-REF.LENGTH)**2

TYPE1	TYPE2	STRETCHK1	REF.LENGTH
1	2	346.0000	1.0890
1	3	331.2000	1.1000
2	2	670.0000	1.3350
2	3	319.5000	1.5010
3	3	316.8000	1.5200

GLASGOW ALKANE/ALKENE FORCE FIELD

PAGE NUMBER 3

VAN DER WAALS

EXPRESSION: EV=((-2.0*EPSILON)/(ALPHA)**6)+(EPSILON*EXP(12.0*(1.0-ALPHA)))

ALPHA=DIST(1-2)/(RSTAR1+RSTAR2)

TYPE1	TYPE2	RSTAR1	RSTAR2	EPSILON
1	1	3.1000	0.0	0.0160
1	2	3.5300	0.0	0.0330
1	3	3.3500	0.0	0.0299
2	2	4.0000	0.0	0.0760
2	3	3.6000	0.0	0.0800
3	3	3.8500	0.0	0.1200

ANGLE BENDING

EXPRESSION: EA=BENDK1*((DELTH)**2)-(BENDK2*(DELTH)**3))

DELTH=ABS(REF.ANGLE"N"-OBSERVED ANGLE)

REF.ANGLE"N" IS USED FOR DEGREE OF NON-HYDROGEN SUBSTITUTION "N" ON THE CENTRAL ATOM OF THE ANGLE CONCERNED. ANGLES ARE IN DEGREES

TYPE1	TYPE2	TYPE3	BENDK1	BENDK2	REF.ANG1	REF.ANG2	REF.ANG3	REF.ANG4
1	2	1	0.0055	0.0096	118.6000	0.0	0.0	0.0
1	2	2	0.0060	0.0	121.7000	120.4000	0.0	0.0
1	2	3	0.0060	0.0	0.0	117.5000	0.0	0.0
2	2	3	0.0120	0.0096	0.0	122.3000	121.0000	0.0
3	2	3	0.0233	0.0096	0.0	0.0	116.4000	0.0
1	3	1	0.0072	0.0096	108.2000	109.1000	0.0	0.0
1	3	2	0.0088	0.0096	110.5000	110.0000	110.2000	0.0
1	3	3	0.0088	0.0096	109.5000	109.0000	109.2000	0.0
2	3	2	0.0090	0.0096	0.0	111.0000	110.1000	109.5000
2	3	3	0.0090	0.0096	0.0	110.4000	110.1000	109.5000
3	3	3	0.0120	0.0096	0.0	110.4000	110.1000	109.5000

GLASGOW ALKANE/ALKENE FORCE FIELD

PAGE NUMBER 5

TORSION

EXPRESSION: $ET=BARRIER\ HEIGHT*(1.0+SIGN*COS(FOLD*W))$

"RANGE" ALLOWS CALCULATION OF TORSIONAL ENERGY ONLY WHEN TORSION ANGLE IS WITHIN THE SPECIFIED RANGE
"W" IS THE TORSION ANGLE

TYPE1	TYPE2	TYPE3	TYPE4	BARRIER HT	SIGN	FOLD	RANGE(+--)
1	2	2	1	6.2500	-1.0	2.0	0.0
1	2	2	3	6.2500	-1.0	2.0	0.0
3	2	2	3	6.2500	-1.0	2.0	0.0
1	2	3	1	0.1367	1.0	3.0	0.0
1	2	3	2	0.1367	1.0	3.0	0.0
1	2	3	3	0.1367	1.0	3.0	0.0
2	2	3	1	0.0629	-1.0	3.0	0.0
2	2	3	2	0.0629	-1.0	3.0	0.0
2	2	3	3	0.0629	-1.0	3.0	0.0
3	2	3	1	0.0629	1.0	3.0	0.0
3	2	3	2	0.0629	1.0	3.0	0.0
3	2	3	3	0.0629	1.0	3.0	0.0
1	3	3	1	0.1100	1.0	3.0	0.0
1	3	3	2	0.1100	1.0	3.0	0.0
1	3	3	3	0.1100	1.0	3.0	0.0
2	3	3	2	0.0200	1.0	3.0	0.0
2	3	3	3	0.0629	1.0	3.0	0.0
3	3	3	3	0.0629	1.0	3.0	0.0

GLASGOW ALKANE/ALKENE FORCE FIELD

PAGE NUMBER 6

OUT OF PLANE BENDING

EXPRESSION: $EO=OP-BENDK*(180.0-CHI)**2$

WHERE "CHI" IS THE IMPROPER TORSION ANGLE DEFINED BY WARSHEL, LEVITT, AND LIFSON
(J.MOL.SPECTROSCOPY, 1970, ~~33~~, 84)

TYPE1	TYPE2	TYPE3	TYPE4	OP-BENDK
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3	2	2	1	0.0020
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APPENDIX B

FFII ALKANE/ALKENE FORCE FIELD (ALLINGER BASED)

ATOM KEY

1 = C(SP)**2, 2 = C(SP)**3, 3 = BR, 4 = H,

FORCE CONSTANTS ARE IN KCAL/MOLE/ANGSTROM**2 OR IN KCAL/MOLE/DEGREE**2
ENERGIES ARE IN KCAL/MOLE

ALKANE/ALKENE/ALKANE FORCE FIELD (ALLINGER BASED) NUMBER PAGE NUMBER 2

NEW DELA WAA BOND STRETCHING

EXPRESSION: EB=STRETCHK1*(LENGTH-REF.LENGTH)**2 PHA/(0.0750)*EPSILON)

ALPHA=VIST(1-2)/((BSTAT1+VSTAT2)

TYPE1	TYPE2	STRETCHK1	REF.LENGTH
1	1	690.8640	1.3320
1	2	316.6460	1.4960
1	4	331.0390	1.0900
2	2	316.6460	1.5300
2	3	179.7200	1.9300
2	4	331.0000	1.0940
1	2	1.5500	0.1160
1	3	1.5500	0.10.2460
1	4	1.5500	0.10.0840
2	3	1.5500	0.5220
2	4	1.5500	0.1770
2	4	1.5500	0.0600

ALKANE/ALKENE FORCE FIELD (ALLINGER BASED)

ALPHA=ALPHA/180

VAN DER WAALS INTERACTIONS

EXPRESSION: $EV = ((-2.25 / (\alpha * 6)) * \epsilon) + (828000 * \exp(-\alpha * 0.0736) * \epsilon)$

ALPHA=DIST(1-2)/(RSTAR1+RSTAR2)

TYPE1	TYPE2	TYPE3	REF-ANG1	RSTAR1	REF-ANG2	RSTAR2	REF-ANG3	EPSILON
1	1	1	0.0000	1.8500	0.0000	1.8500	0.0000	0.0330
1	1	2	0.0053	1.8500	0.0000	1.5500	0.0000	0.0620
1	1	3	0.0088	1.8500	0.0000	1.9500	0.0000	0.0440
1	1	4	0.0088	1.8500	0.0000	1.4500	0.0000	0.0440
2	2	2	0.0053	1.5500	0.0000	1.5500	0.0000	0.1160
2	2	3	0.0088	1.5500	0.0000	1.9500	0.0000	0.2460
2	2	4	0.0088	1.5500	0.0000	1.4500	0.0000	0.0840
3	3	3	0.0053	1.9500	0.0000	1.9500	0.0000	0.5220
3	3	4	0.0053	1.9500	0.0000	1.4500	0.0000	0.1770
4	4	4	0.0045	1.4500	0.0000	1.4500	0.0000	0.0600

ALKANE/ALKENE FORCE FIELD (ALLINGER BASED)

PAGE NUMBER 4

ANGLE BENDING

EXPRESSION: EA=BENDK1*(ANGLE-REF.ANGLE"N")**2

"N" REFERS TO DEGREE OF NON-HYDROGEN SUBSTITUTION ON THE CENTRAL ATOM OF THE ANGLE CONCERNED. ANGLES ARE IN DEGREES

TYPE1	TYPE2	TYPE3	BENDK1	REF.ANG1	REF.ANG2	REF.ANG3	REF.ANG4
1	1	2	0.0088	0.0	122.2000	0.0	0.0
1	1	4	0.0053	0.0	119.9000	0.0	0.0
2	1	4	0.0053	0.0	117.9000	0.0	0.0
1	2	2	0.0088	0.0	110.2000	110.6000	0.0
1	2	3	0.0088	0.0	0.0	110.0000	0.0
1	2	4	0.0053	0.0	112.8000	108.4000	0.0
2	2	2	0.0088	0.0	110.2000	110.6000	109.5000
2	2	3	0.0088	0.0	110.0000	110.0000	109.5000
2	2	4	0.0053	107.8000	112.8000	108.4000	0.0
3	2	4	0.0053	0.0	109.0000	109.0000	0.0
4	2	4	0.0044	111.2000	108.5000	0.0	0.0

ALKANE/ALKENE FORCE FIELD (ALLINGER BASED) PAGE NUMBER 5

TORSION

EXPRESSION: $EI = \text{BARRIER HEIGHT} * (1.0 + \text{SIGN} * \cos(\text{FOLD} * W))$

"RANGE" ALLOWS CALCULATION OF TORSIONAL ENERGY ONLY WHEN TORSION ANGLE IS WITHIN THE SPECIFIED RANGE
"W" IS THE TORSION ANGLE

TYPE1	TYPE2	TYPE3	TYPE4	BARRIER HT	SIGN	FOLD	RANGE (+--)
2	1	1	2	6.7700	-1.0	2.0	0.0
2	1	1	4	6.7700	-1.0	2.0	0.0
4	1	1	4	6.7700	-1.0	2.0	0.0
1	1	2	2	0.1541	-1.0	3.0	0.0
1	1	2	3	0.1541	-1.0	3.0	0.0
1	1	2	4	0.1541	-1.0	3.0	0.0
4	1	2	2	0.1541	1.0	3.0	0.0
4	1	2	3	0.1541	1.0	3.0	0.0
4	1	2	4	0.1541	1.0	3.0	0.0
1	2	2	1	0.1000	1.0	3.0	0.0
1	2	2	2	0.1000	1.0	3.0	0.0
1	2	2	3	0.1000	1.0	3.0	0.0
1	2	2	4	0.1000	1.0	3.0	0.0
2	2	2	4	0.1000	1.0	3.0	0.0
3	2	2	4	0.1000	1.0	3.0	0.0
4	2	2	4	0.1000	1.0	3.0	0.0

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ALKANE/ALKENE FORCE FIELD (ALLINGER BASED)

OUT OF PLANE BENDING

EXPRESSION: $EO=OP-BENDK*(180.0-CHI)**2$

WHERE "CHI" IS THE IMPROPER TORSION ANGLE DEFINED BY WARSHEL, LEVITT, AND LIFSON (J. MOL. SPECTROSCOPY, 1970, ~~33~~, 84)

TYPE1	TYPE2	TYPE3	TYPE4	OP-BENDK
3	2	2	1	0.0035

APPENDIX C

PAGE NUMBER 2

SESQUITERPENE FORCE FIELD

BOND STRETCHING

FFIII SESQUITERPENE FORCE FIELD

ATOM	KEY	TYPE1	TYPE2	STRETCHING1	REF. LENGTH
1	= C(SP)**2, 2 = C(=O)			3 = C(SP)**3, 4 = O(=C)	5 = -O-
6	= H				
		1	3	316.8460	1.4960
		1	9	331.0390	1.0900
		2	3	316.8460	1.4960
		2	4	777.2220	1.2100
		2	5	674.8648	1.3340
		3	3	316.8460	1.4960
		3	9	331.0390	1.0940
		5	3	503.7500	0.9520

FORCE CONSTANTS ARE IN KCAL/MOLE/ANGSTROM**2 OR IN KCAL/MOLE/DEGREE**2
ENERGIES ARE IN KCAL/MOLE

SE50 SESQUITERPENE FORCE FIELD

PAGE NUMBER 2

VAN BOND STRETCHING

EXPRESSION: EB=STRETCHK1*(LENGTH-REF.LENGTH)**2

TYPE1	TYPE2	STRETCHK1	REF.LENGTH
1	1	690.8640	1.3320
1	2	316.6460	1.4800
1	3	316.6460	1.4960
1	6	331.0390	1.0900
2	3	316.6460	1.4960
2	4	777.2220	1.2100
2	5	474.9690	1.3340
3	3	316.6460	1.5300
3	5	309.4500	1.4700
3	6	331.0390	1.0940
5	6	503.7500	0.9520

SESQUITERPENE FORCE FIELD

PAGE NUMBER 4

ANGLE BENDING

EXPRESSION: EA=BENDK1*(ANGLE-REF.ANGLE"UN")**2

"UN" REFERS TO DEGREE OF NON-HYDROGEN SUBSTITUTION ON THE CENTRAL ATOM OF THE ANGLE CONCERNED. ANGLES ARE IN DEGREES

TYPE1	TYPE2	TYPE3	BENDK1	REF.ANG1	REF.ANG2	REF.ANG3	REF.ANG4
1	1	2	0.0088	0.0	122.2000	121.7000	0.0
1	1	3	0.0088	0.0	122.2000	121.7000	0.0
1	1	6	0.0053	118.9000	119.9000	0.0	0.0
2	1	3	0.0088	0.0	0.0	116.6000	0.0
2	1	6	0.0053	0.0	117.9000	0.0	0.0
3	1	3	0.0088	0.0	0.0	116.6000	0.0
3	1	6	0.0053	0.0	117.9000	0.0	0.0
6	1	6	0.0044	122.2000	0.0	0.0	0.0
1	2	1	0.0088	0.0	0.0	115.6000	0.0
1	2	4	0.0125	0.0	122.2000	122.2000	0.0
1	2	5	0.0123	0.0	0.0	115.6000	0.0
3	2	4	0.0125	0.0	124.4000	124.4000	0.0
3	2	5	0.0123	0.0	0.0	115.6000	0.0
4	2	5	0.0156	0.0	0.0	120.0000	0.0
1	3	1	0.0088	0.0	110.2000	110.6000	109.5000
1	3	3	0.0088	0.0	110.2000	110.6000	109.5000
1	3	5	0.0110	0.0	110.2000	110.6000	109.5000
1	3	6	0.0053	107.8000	112.8000	108.4000	0.0
2	3	3	0.0088	0.0	110.2000	110.6000	109.5000
2	3	6	0.0053	107.8000	108.5000	108.4000	0.0
3	3	3	0.0088	0.0	110.2000	110.6000	109.5000
3	3	5	0.0110	0.0	110.2000	110.6000	109.5000
3	3	6	0.0053	107.8000	112.8000	108.4000	0.0
5	3	6	0.0097	107.8000	112.8000	108.4000	0.0
6	3	6	0.0044	111.2000	108.5000	0.0	0.0
2	5	3	0.0221	0.0	114.8000	0.0	0.0
3	5	6	0.0167	95.2000	0.0	0.0	0.0

SESQUITERPENE FORCE FIELD

PAGE NUMBER 5

TORSION

EXPRESSION: $ET=BARRIER\ HEIGHT*(1.0+SIGN*COS(FOLD*W))$

"RANGE" ALLOWS CALCULATION OF TORSIONAL ENERGY ONLY WHEN TORSION ANGLE IS WITHIN THE SPECIFIED RANGE
"W" IS THE TORSION ANGLE

TYPE1	TYPE2	TYPE3	TYPE4	BARRIER HT	SIGN	FOLD	RANGE (+--)
2	1	1	3	6.7700	-1.0	2.0	0.0
3	1	1	3	6.7700	-1.0	2.0	0.0
2	1	1	6	6.7700	-1.0	2.0	0.0
3	1	1	6	6.7700	-1.0	2.0	0.0
6	1	1	6	6.7700	-1.0	2.0	0.0
1	1	2	1	1.3000	-1.0	2.0	0.0
1	1	2	4	1.3000	-1.0	2.0	0.0
1	1	2	5	1.3000	-1.0	2.0	0.0
3	1	2	1	1.3000	-1.0	2.0	0.0
3	1	2	4	1.3000	-1.0	2.0	0.0
3	1	2	5	1.3000	-1.0	2.0	0.0
6	1	3	1	0.1800	-1.0	3.0	0.0
1	1	3	3	0.1800	-1.0	3.0	0.0
1	1	3	5	0.1800	-1.0	3.0	0.0
1	1	3	6	0.1800	-1.0	3.0	0.0
2	1	3	3	0.1800	1.0	3.0	0.0
2	1	3	6	0.1800	1.0	3.0	0.0
3	1	3	1	0.1800	1.0	3.0	0.0
3	1	3	5	0.1800	1.0	3.0	0.0
6	1	3	6	0.1800	1.0	3.0	0.0
6	1	3	1	0.1600	1.0	3.0	0.0

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SESQUITERPENE FORCE FIELD

TORSION (CONTINUED)

TYPE1	TYPE2	TYPE3	TYPE4	BARRIER HT	SIGN	FOLD	RANGE (+--)
6	1	3	3	0.1600	1.0	3.0	0.0
6	1	3	5	0.1800	1.0	3.0	0.0
6	1	3	6	0.1800	1.0	3.0	0.0
4	2	3	3	-0.7450	1.0	3.0	60.0
4	2	3	6	-0.3050	1.0	3.0	60.0
5	2	3	3	0.0450	1.0	3.0	0.0
5	2	3	6	-0.0200	1.0	3.0	0.0
1	2	5	3	5.9210	-1.0	2.0	0.0
3	2	5	3	5.9210	-1.0	2.0	0.0
4	2	5	3	5.9210	-1.0	2.0	0.0
1	3	3	1	0.0830	1.0	3.0	0.0
1	3	3	3	0.0830	1.0	3.0	0.0
1	3	3	5	0.0830	1.0	3.0	0.0
1	3	3	6	0.0830	1.0	3.0	0.0
2	3	3	3	0.0830	1.0	3.0	0.0
3	3	3	3	0.0830	1.0	3.0	0.0
5	3	3	3	0.0830	1.0	3.0	0.0
2	3	3	6	0.0830	1.0	3.0	0.0
3	3	3	6	0.0830	1.0	3.0	0.0
5	3	3	6	0.0830	1.0	3.0	0.0
6	3	3	6	0.0830	1.0	3.0	0.0
1	3	5	2	0.2000	1.0	3.0	0.0
1	3	5	6	0.2000	1.0	3.0	0.0
3	3	5	2	0.2000	1.0	3.0	0.0
3	3	5	6	0.2000	1.0	3.0	0.0
6	3	5	2	0.2000	1.0	3.0	0.0
6	3	5	6	0.2000	1.0	3.0	0.0

SESQUITERPENE FORCE FIELD

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OUT OF PLANE BENDING

EXPRESSION: $E0=OP-BENDK*(180.0-CHI)**2$

WHERE "CHI" IS THE IMPROPER TORSION ANGLE DEFINED BY WARSHEL,LEVITT,AND LIFSON
(J.MOL.SPECTROSCOPY,1970,33,84)

TYPE1	TYPE2	TYPE3	TYPE4	OP-BENDK
3	1	1	6	0.0035
3	1	1	3	0.0035
5	2	1	4	0.0092
5	2	3	4	0.0092

